more here a 187 of 197 of 198 of 198 of 198 of 198 of 198 of 199 of 199 AN 22270, AI190178 AN AN 38936 AN A				- 1					
HLYDIO4 876169 Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 487 of SEQ ID NO:1378, b is an integer of 15 to 501, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1378, and where b is greater than or equal to a + 14. HBXFF23 876170 Preferably excluded from the polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 948 of SEQ ID NO:1379, b is an integer of 15 to 962, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1379, and where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1379, and where b is greater than or equal to a + 14. HDPBG07 876172 Preferably excluded from the polynucleotides comprising a polynucleotides comprising a polynucleotide sequence described by the general formula of a-b, where a list any integer between 1 to 2921 of A150895 H160518 is any integer between 1 to 2921 of A160852 W68464, SEQ ID NO:1380, b is an integer of 15 to 2935, where both a and b No:20395, A163995, A165995				(
HLYDIO4 876169 Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 487 of 520 ID No:1378, b is an integer of 15 to 501, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1378, and where b is greater than or equal to a + 14. HBXFF23 876170 Preferably excluded from the present invention are one or more polynucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 948 of 5EQ ID NO:1379, b is an integer of 15 to 962, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1379, and where b is greater than or equal to a + 14. HDPBG07 876172 Preferably excluded from the present invention are one or more A1550191, AA202250, HIGHOS18 is any integer between 1 to 2921 of A113168, AA062951 is any integer between 1 to 2921 of A113168, AA062951 is any integer between 1 to 2921 of A113655, A135995, A1555951				than or equal to a + 14.					
present invention are one or more present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 487 of SEQ ID NO:1378, b is an integer of nucleotide residues shown in SEQ ID NO:1378, and where b is greater than or equal to a + 14. HBXFF23 876170 Preferably excluded from the polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 948 of SEQ ID NO:1379, b is an integer of 15 to 962, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1379, and where b is greater than or equal to a + 14. HDPBG07 876172 Preferably excluded from the AM450363, AA8605227 preferably excluded from the AM450363, AA8605227 preferably excluded from the AM450363, AA860528 is any integer between 1 to 2921 of A113168, AA662911 polynucleotide sequence described by AN138656, A1630596 correspond to the positions of A1138656, A1630596	1378	HLYDI04	876169	Preferably excluded from the					
polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 487 of SEQ ID NO:1378, b is an integer of 15 to 501, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1378, and where b is greater than or equal to a + 14. HBXFF23 876170 Preferably excluded from the polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a nucleotide sequence described by the general formula of a-b, where a sex of is any integer between 1 to 948 of SEQ ID NO:1379, b is an integer of 15 to 962, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1379, and where b is greater than or equal to a + 14. HDPBG07 876172 Preferably excluded from the present invention are one or more Al191168, AA806221 polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a NISC1919, AA306231 is any integer between 1 to 2921 of AI030536,				present invention are one or more					
the general formula of a-b, where a is any integer between the positions of nucleotide residues shown in SEQ ID No.1378, b is an integer of 15 to 501, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO.1378, and where b is greater than or equal to a + 14. HBXFF23 876170 Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 948 of SEQ ID NO.1379, b is an integer of 15 to 962, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO.1379, and where b is greater than or equal to a + 14. HDPBG07 876172 Preferably excluded from the polynucleotide residues shown in SEQ ID NO.1379, and where b is greater than or equal to a + 14. HDPBG07 876172 Preferably excluded from the polynucleotide sequence described by preferably excluded from the polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a AIS21191, AA20231 the general formula of a-b, where a lis any integer between 1 to 2921 of AI138636, AI143991 correspond to the positions of AI139995, AIS65951				polynucleotides comprising a					
the general formula of a-b, where a is any integer between 1 to 487 of SEQ IN 0:1378, b is an integer of 15 to 501, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1378, and where b is greater than or equal to a + 14. HBXFF23 876170 Preferably excluded from the present invention are one or more polymucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 948 of SEQ ID NO:1379, b is an integer of 15 to 962, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1379, and where b is greater than or equal to a + 14. HDPBG07 876172 Preferably excluded from the AM450363, AA806222 present invention are one or more A113168, AA062917 preferably excluded from the A113168, AA062917 in any integer between 1 to 2921 of A1130178 A1190178 is any integer between 1 to 2921 of A1160853 (SEQ ID NO:1380, b is an integer of A123991 correspond to the positions of A123995, A1565861				nucleotide sequence described by					-
is any integer between 1 to 487 of SEQ ID NO:1378, b is an integer of 15 to 501, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1378, and where b is greater than or equal to a + 14. HBXFF23 876170 Preferably excluded from the polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 948 of SEQ ID NO:1379, b is an integer of 15 to 962, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1379, and where b is greater than or equal to a + 14. HDPBG07 876172 Preferably excluded from the polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a AIH3168, AA062917 nucleotide sequence described by the general formula of a-b, where a AIH3168, AA062917 sea my integer between 1 to 2921 of AIN316935, AIE60852 15 to 2935, where both a and b AIN3636, AIA33991 correspond to the positions of AIN3636, AIA33991				ø					
SEQ ID NO:1378, b is an integer of 15 to 501, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1378, and where b is greater than or equal to a + 14. HBXFF23 876170 Preferably excluded from the polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 948 of SEQ ID NO:1379, b is an integer of 15 to 962, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1379, and where b is greater than or equal to a + 14. HDPBG07 876172 Preferably excluded from the polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a AI143168, AA062917 nucleotide sequence described by the general formula of a-b, where a AI143168, AA062917 secont invention are one or more polynucleotides comprising a AI143168, AA06291 is any integer between 1 to 2921 of AN13636, AI160536 is any integer between 1 to 2921 of AN13636, AI1433991 correspond to the positions of AN13636, AI433991				ny integer between 1 to 487					
15 to 501, where both a and b Correspond to the positions of NO:1378, and where b is greater than or equal to a + 14. HBXFF23 876170 Preferably excluded from the polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 948 of SEQ ID NO:1379, b is an integer of 15 to 962, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1379, and where b is greater than or equal to a + 14. HDPBG07 876172 Preferably excluded from the polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a AIH3168, AA062917 nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 2921 of AIT50852, We8464, SEQ ID NO:1380, b is an integer of AN138636, AIE50595 15 to 2935, where both a and b AIT779995, AIS6595									
correspond to the positions of nucleotide residues shown in SEQ ID NO:1378, and where b is greater than or equal to a + 14. HBXFF23 876170 Preferably excluded from the polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 948 of SEQ ID NO:1379, b is an integer of 15 to 962, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1379, and where b is greater than or equal to a + 14. HDPBG07 876172 Preferably excluded from the polynucleotides comprising a AM450363, AA806222 present invention are one or more polynucleotides comprising a AM450363, AA806222 present invention are one or more polynucleotides comprising a AM450363, AA806222 present invention are one or more polynucleotides comprising a AM450363, AA806222 present invention are one or more polynucleotides comprising a AM450363, AA806222 present invention are one or more polynucleotides comprising a AM450363, AA806222 present invention are one or more polynucleotides comprising a AM450363, AA806222 present invention are one or more polynucleotides comprising a AM450363, AA806222 present invention are one or more polynucleotides comprising a AM450363, AA806222 present invention are one or more polynucleotides comprising a AM450363, AA806222 present invention are one or more polynucleotides comprising a AM450363, AA806222 present invention are one or more polynucleotides comprising a AM450363, AA806222 present invention are one or more polynucleotide sequence described by AM450363, AA806222 present invention are one or more polynucleotides comprising a AM450363, AA806222 present invention are one or more polynucleotides comprising a AM450363, AA806222 present invention are one or more polynucleotide sequence of a more				15 to 501, where both a and b					
NO:1378, and where b is greater Than or equal to a + 14. HBXFF23 876170 Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 948 of 15 to 962, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1379, and where b is greater than or equal to a + 14. HDPBG07 876172 Preferably excluded from the polynucleotides comprising a nucleotide sequence described by AM450363, A88062270, A1190178 polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a HI350341, AA477713 polynucleotide sequence described by AM722270, A1190178 is any integer between 1 to 2921 of A1160536 15 to 2935, where both a and b A1279995, A156596				correspond to the positions of					
HDRSFF23 876170 Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 948 of SEQ ID NO:1379, b is an integer of 15 to 962, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1379, and where b is greater than or equal to a + 14. HDPBG07 876172 Preferably excluded from the present invention are one or more polynucleotides comprising a present invention are one or more polynucleotides comprising a A1524191, AA280231 the general formula of a-b, where a A154191, AA280231 is any integer between 1 to 2921 of A11808363, A160852 is any integer between 1 to 2921 of A11808363, A1828995 a156596		_		Ö					
HBXFF23 876170 Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 948 of 15 o 962, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1379, and where b is greater than or equal to a + 14. HDPBG07 876172 Preferably excluded from the polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a nucleotide sequence described by the general formula of a-b, where a list any integer between 1 to 2921 of AI1938636, AI160852 (SEQ ID NO:1380, b is an integer of AI138636, AI163399] correspond to the positions of AI279995, AI56596				and where b is					
HBXFF23 876170 Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 948 of 15 to 962, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1379, and where b is greater than or equal to a + 14. HDPBG07 876172 Preferably excluded from the polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a nucleotide sequence described by the general formula of a-b, where a sis any integer between 1 to 2921 of AI760852, W68464, SEQ ID NO:1380, b is an integer of AI133836, AI1433991 correspond to the positions of AI279995, AI56596				equal to a + 14.					
present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 948 of SEQ ID NO:1379, b is an integer of 15 to 962, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1379, and where b is greater than or equal to a + 14. HDPBG07 876172 Preferably excluded from the polynucleotides comprising a AM450363, AA806222 present invention are one or more polynucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 2921 of SEQ ID NO:1380, b is an integer of AI19186, AI160536 correspond to the positions of AI279995, AI56596	1379	HBXFF23	876170	Preferably excluded from the	W03002				
polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 948 of SEQ ID NO:1379, b is an integer of 15 to 962, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1379, and where b is greater than or equal to a + 14. HDPBG07 876172 Preferably excluded from the polynucleotides comprising a AA722270, A1191078 the general formula of a-b, where a A1143168, AA662917 nucleotide sequence described by the general formula of a-b, where a A1524191, AA280235 is any integer between 1 to 2921 of A1760852, W68464, SEQ ID NO:1380, b is an integer of AN138636, A14239995, A1565966				present invention are one or more					
the general formula of a-b, where a is any integer between 1 to 948 of SEQ ID NO:1379, b is an integer of 15 to 962, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1379, and where b is greater than or equal to a + 14. HDPBG07 876172 Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by AA722270, AI190176 the general formula of a-b, where a AI524191, AA280235 is any integer between 1 to 2921 of AI760852, W68464, SEQ ID NO:1380, b is an integer of AN38636, AI423991 correspond to the positions of AI279995, AIS65961				polynucleotides comprising a					
the general formula of a-b, where a is any integer between 1 to 948 of SEQ ID NO:1379, b is an integer of 15 to 962, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1379, and where b is greater than or equal to a + 14. HDPBG07 876172 Preferably excluded from the polynucleotides comprising a AN450363, AA806221 polynucleotides comprising a AN722270, AI1910178 the general formula of a-b, where a AI524191, AA280238 is any integer between 1 to 2921 of AI760852, W68464, SEQ ID NO:1380, b is an integer of AU38636, AI423991 correspond to the positions of AI279995, AIS65961				nucleotide sequence described by					
is any integer between 1 to 948 of SEQ ID NO:1379, b is an integer of 15 to 962, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1379, and where b is greater than or equal to a + 14. HDPBG07 876172 Preferably excluded from the present invention are one or more present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a AI143168, AA062917 nucleotide sequence described by the general formula of a-b, where a AI524191, AA280239 is any integer between 1 to 2921 of AI760852, W68464, SEQ ID NO:1380, b is an integer of AI097247, AI160536 correspond to the positions of AI279995, AI565961		-		the general formula of a-b, where a					
SEQ ID NO:1379, b is an integer of 15 to 962, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1379, and where b is greater than or equal to a + 14. HDPBG07 876172 Preferably excluded from the polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 2921 of SEQ ID NO:1380, b is an integer of 15 to 2935, where both a and b AN138636, AI423991 correspond to the positions of A1779995, AI565961				ny integer between 1 to 948					
15 to 962, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1379, and where b is greater than or equal to a + 14. HDPBG07 876172 Preferably excluded from the polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 2921 of SEQ ID NO:1380, b is an integer of AIT316852, M68464, SEQ ID NO:1380, b is an integer of AIT316853, AIT33991 AM138636, AIT33991 AIT3169536									
correspond to the positions of nucleotide residues shown in SEQ ID NO:1379, and where b is greater than or equal to a + 14. HDPBG07 876172 Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 2921 of SEQ ID NO:1380, b is an integer of AN138636, AN123991 correspond to the positions of AN38636, AN565961	•			15 to 962, where both a and b					
HDPBG07 876172 Preferably excluded from the positions of Alson are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a sis any integer between 1 to 2921 of AM138636, AI1433991 correspond to the positions of AI279995, AI565961				correspond to the positions of					
than or equal to a + 14. HDPBG07 876172 Preferably excluded from the polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a h14316852, w68464, SEQ ID NO:1380, b is an integer of the positions of AII38636, AII60536 also correspond to the positions of AII38636, AII565961									,
HDPBG07 876172 Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a h1524191, AA280235 is any integer between 1 to 2921 of h160852, W68464, SEQ ID NO:1380, b is an integer of h1760852, M1423991 correspond to the positions of h1279995, AIS65961									
HDPBG07 876172 Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a hstatistical by sEQ ID NO:1380, b is an integer of sequence both a and b halfors AII50536				equal to a +					
A1950341, AA477713 A1143168, AA062917 AA722270, A1190178 A A1524191, AA280235 of A1760852, W68464, A1097247, A1160536 AN138636, A1423991 AI279995, AI565961	1380	HDPBG07	876172	Preferably excluded from the	AW450363,	AA806222,	AI697498, A	AW379227,	
AI143168, AA062917 AA722270, AI190178 A AI524191, AA280235 of AI760852, W68464, AI097247, AI160536 AW138636, AI423991 AI279995, AI565961					AI950341,	AA477713,	_	AI762090,	
a AI524191, AA280235 of AI760852, W68464, aI097247, AI160536 AW138636, AI423991 AI279995, AI565961				polynucleotides comprising a	AI143168,	AA062917,	AW055125, A	AI708563,	
a AIS24191, AA280235 of AI760852, W68464, aI097247, AI160536 AW138636, AI423991 AI279995, AIS65961				nucleotide sequence described by	AA722270,	AI190178,	AI147612, A	AA188072,	
of A1760852, W68464, A1097247, A1160536 AW138636, A1423991 AI279995, AIS65961					AI524191,	AA280235,	N44673, AI921393, AI291105	21393, AI2	291105,
of AI097247, AW138636, AI279995,				is any integer between 1 to 2921 of	AI760852,	W68464, N2	N26444, AI373000, AI302843,	1000, AI302	2843,
AW138636, AI279995,				SEQ ID NO:1380, b is an integer of	AI097247,	AI160536,	T66196, AI8	AI804233, W78	W78020,
of AI279995,				15 to 2935, where both a and b	AW138636,	AI423991,	AI089967, C	C75569, AA	AA565899,
				correspond to the positions of	AI279995,	AIS65961,	AW341212, H	H99338, AI	AI299654,
SEQ ID AA631426,				nucleotide residues shown in SEQ ID	AA631426,	AA419222,	AA663984, W	W73977, AAS	AA954140,

			NO:1380, and where b is greater	W51950, W69512, AA410280, AI491793, AI393820,
			than or equal to a + 14.	
				N35301, N4
				H43944, AW407957, AI186159, N95537, AA730169,
				W73396, H96739, AA0824
				, N33412, C75660, AI35
	_			AW291308, AI656702, AI242486, AW026628,
				T91940, AI693563, R81438, AI868184, H42592,
				AA419207
				H23921, R81641, AA345108, AA361827, AI707909,
				AA310049, AA346697, W69413, AW407592, T66132,
	_			
				i, T10717
				A928401, AI
				AA761812, W69429, AA922945, AI381590, AI347968,
	_			N24768, AR038868, AB016811, AR055261, AR038869,
				AR055262
1381 HCY	HCYBF02 87	876174	Preferably excluded from the	, AA134366, AA259244
			present invention are one or more	1591375,
			prising a	, AW338376,
_			ednence de	7, AL035847, N54947,
	_		the general formula of a-b, where a	N80390, AL039471, AA078337, AA515176, AW008089,
			is any integer between 1 to 612 of	AA171400, AA595499, AW247866, AW250983, T94247,
	_		b is an inte	٦,
			15 to 626, where both a and b	AA833896, AA833875, AA493464, AW168520,
			correspond to the positions of	AA350593, AA610381, AA568494, AI952885,
			$\boldsymbol{\tau}$	AA772140, AL044674, AW080062, AA526542,
			NO:1381, and where b is greater	AA847711, AA665645, AA601674, AA335314,
			than or equal to a + 14.	, AI050050, AW088745,
		1		AI224583, AA320262, AA847095, AA493136,
				AI064918, AA743517, AI000381, AA595661, N59648,

AA679937, AW029515, AA666052,	AA218684, AA548390, AA584862, AA283455, AI440037, AA613627, AA524604, AI583321, F31380,	AL118823, AA199578, AW021105,	661, F18553, AW419209,	, AA551105, R92608	, H88429, AI927275,	AA687730, AA634882, H62123, AW169038, AA071173,	AL022330,	AC004914, Z77249, AC004973, AF196970, AL079339,	w	AP000031, Z75744, AL031293, AC006539, AC003668,	AC005549, AL121578, AL049636, L05367, AP000038,	AC004929,	AL035086, AC000115, AL031283, AL022165,	AC004526, AP000135,	, AP000106, AC007308	Z95125, AL035413, AC006251, AL109865, AL031073,		0, AF165141, AC006509, AC005484	Z98049, AC011456,	AC004079,	9, AC004263, AC004988, AL035653	, U91326, AC005412, AC002425,	, AL009047, AC007533, Z83826, AC	, AF207955, AL035460	, L29074, AP000261, AF	AL031652, AC005632, AC007463, AC005209,	AC002403, AP000100, AP000035, AC005048,	٥,	AL022322, AC006241, AL080239, AC002395,	9, AL133396	, Z69705, AC004063, AL	5960, AJ131016, AC004754,	AC005046, AC002110, AJ006345, AC005832,
				•																				-									

		AC005829, AP000010	010, AC004961, AC005725,	
		AL022239, AC00210	5, Z98050, AC005225	AC006270.
		-	U82828, AC008064,	AP000247,
		AC007066, AP000255	, AL049832, Z84484,	284572,
		AC004853, AC002039	, AC006062,	
		AL031733, AP000497,	Z97353, AP000503,	AL133353,
		AL008712, AC005377	_	
		AC005690, AC004938	938, AC007388, AC005876,	
		AC006142, AP000102	, AL034429	-
		AL121576, AC002492	, Z73358, AP000351,	AC008372,
		AC009399, Z97184,	AL049829, AC004099,	AC007538,
			, AL122003	
		AP000201, AC007539	539, AL022328, AF049895,	
		AC002064, AC006385	385, AC005042, AC007955,	
		AC007731, AC004975	975, AP000097, AC007682,	-,-
		AL049712, AL022163	163, AC009248, AL031985,	
		AC006155, AP000356	356, AC005191, AC006965,	
-		AC007385, AC005988	_	
		AC005409, AL023095	, AC004953,	
			, Z84469, AC005500,	AL021331,
			, AF165926, Y10196,	AP000354,
			_	
			5, U82696, AP000338,	AL132987,
		U	AC007200, AP000216,	AC003098,
			, AC005342,	
		_	, AC004815,	
			055, AC004876, AL031729, Z	68287
1382 HTWD[21 876177 Pr	'n		_	
ıd	present invention are one or more		, AA644542,	
od		_	8, AI247781, AI076324,	N68357,
nu	nucleotide sequence described by	AI380870, T87807,	AA808229, AW197425,	AA835077,
the	general formula of a-b,	Z40387, AI458836	9	-
is	teger between 1 to 569			
SE	SEQ ID NO:1382, b is an integer of			
15	where both a and			
00	correspond to the positions of	-		

			nucleotide residues shown in SEQ ID				
			NO:1382, and where b is greater than or equal to a + 14.				
1383	HATED01	876179	Preferably excluded from the	AI792782,	AI191919,	AI765864,	AI733139,
			present invention are one or more	AA702347,	AI220405,	AI423312,	AI478373,
			polynucleotides comprising a	AW302194,	AI423507,	AI916231,	AI627973,
			nucleotide sequence described by	AW173486,	AI086574,	AI701146,	AI521715,
		_	the general formula of a-b, where a	AI917438,	AI678790,	AI925944,	AI770081,
			is any integer between 1 to 503 of	AA760715,	AI904742,	AI582603,	AI990352,
			SEQ ID NO:1383, b is an integer of	AI951007,	AI655622,	AI650463,	AW173518,
			15 to 517, where both a and b	AI393071,	AW236096,	AI989921,	AI022200,
			correspond to the positions of	AI024409,	AI393059,	AI695050,	AA888360,
			nucleotide residues shown in SEQ ID	AI206995,	AI077536,	AI474034,	AI452440,
			NO:1383, and where b is greater	AW194978,	AI076106,	AI206908,	AA969379;
			than or equal to a + 14.	AASS1593,	AI223442,	AI302211,	AI968178,
				AIS71592,	AI241002,	AL034553,	D86198, AF007875,
				AB004789			
1384	HWLVU14	876182	Preferably excluded from the	AI347147,	AI738411,	AI439130,	AA514394,
			present invention are one or more	AA595253,	AI269359,	AW028586,	AI936898,
			_	AI739648,	AW242697,	AW027766,	AA081901,
			nucleotide sequence described by	AI739639,	AW157368,	AI739255,	AI393079,
			the general formula of a-b, where a	AI244459,	AA226866,	N99765, AV	AW418654, AA480225,
			is any integer between 1 to 1216 of	AA905814,	AA999828,	AC007501,	U80736
			SEQ ID NO:1384, b is an integer of				
			15 to 1230, where both a and b				
			correspond to the positions of				
			nucleotide residues shown in SEQ ID				
			NO:1384, and where b is greater				
			than or equal to a + 14.				
1385	HOVCI12	876183	Preferably excluded from the	AA307780,	AI923248		
			present invention are one or more				
			polynucleotides comprising a				
			nucleotide sequence described by				
			the general formula of a-b, where a				
			is any integer between 1 to 368 of				
			SEQ ID NO:1385, b is an integer of				

			15 to 382, where both a and b					
			nucleotide residues shown in SEQ ID					
			NO:1385, and where b is greater					
			than or equal to a + 14.					
1386	HCYBB01	876184	Preferably excluded from the	AW188031,	AI922934,	AA504414,	AI536863,	
			present invention are one or more	AA744849,	AA972022,	AA309130,	AI569395,	
			polynucleotides comprising a	AA135144,	AI570856,	AW021626,	_	
			nucleotide sequence described by	AA962329,	AA737604,	AI351478,	AI560610,	AA765375
			the general formula of a-b, where a					
			is any integer between 1 to 1188 of					
			SEQ ID NO:1386, b is an integer of					
			correspond to the positions of					
			nucleotide residues shown in SEQ ID					
			NO:1386, and where b is greater					
			w					
1387	HCRPM32	876187		AA019767,	AA213771,	H86330, H6	H85652, H867	75,
			present invention are one or more	H86333, AI990107	1990107			
			polynucleotides comprising a					
			nucleotide sequence described by					
			the general formula of a-b, where a					
			· L					
			and the world because the control of					
		-	e E					
			Is to s/s, where both a and b					
			nucleotide residues shown in SEQ ID					
			NO:1387, and where b is greater					_
			than or equal to a + 14.					
1388	HLDNV31	876192	Preferably excluded from the	AI741793,	AW003635,	AA425065,	AL044729,	
			present invention are one or more	AI825212,	AI333124,	AW102958,	AA699738,	
			polynucleotides comprising a	AW014983,	AI580520,	AA653341,	AI248768,	
			nucleotide sequence described by	AW057987,	AA961070,	H11570, A	AA913775, AI	AI425117,
			the general formula of a-b, where a	AI452997,	AI937807,	AL039909,	AL039909, AL041387,	
			is any integer between 1 to 1658 of	AA398627,	AI223186,	T87214, AL045603,		AI638724,
			SEO ID NO:1388, b is an integer of	AA644230,	R45377, A	AI700094, T	T74013, Z21364	64,

				H
			15 to 1672, where both a and b correspond to the positions of	AA/49051, F10219, K14519, A1242930, K40555, R21286, F12602, AA887964, H11462, AA416562,
			nucleotide residues shown in SEQ ID	Z21365, AI890224, R41179, AA829590, AA417298,
			NO:1388, and where b is greater	AA653411, AA837654, AI221436, AA493103,
			than or equal to a + 14.	AW082244, R14339, AA055888, AW389658, T67466, T97917, R08296, AB002326
1389	HCRNN03	876193	Preferably excluded from the	AC005219
			present invention are one or more	
			polynucleotides comprising a	
			nucleotide sequence described by	
			the general formula of a-b, where a	
			is any integer between 1 to 434 of	
			SEQ ID NO:1389, b is an integer of	
			15 to 448, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1389, and where b is greater	
			egual to a + 14.	
1390	HTPIO89	876198		AI808815, AI457550, AI911077, AI658931,
2			present invention are one or more	AI916359, AW009684, AW072228, AA579578,
			polynucleotides comprising a	AA622141, AA295027, AA552628, AA594836,
			nucleotide sequence described by	AA576815,
				AI.127668
			the general lormula of a.b, where a	
		_		
			SEQ ID NO:1390, b is an integer of	
			15 to 882, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1390, and where b is greater	
			than or equal to a + 14.	
1391	HWLQD01	876200	Preferably excluded from the	
			present invention are one or more	
			ŏ	
			nucleotide sequence described by	
			the general formula of a-b, where a	
			is any integer between 1 to 409 of	

er of	SEQ ID	ıe	The state of the s		more		by	nere a	842 of	er of			SEQ ID	J.S.		N24236, AI742828	more		by.	nere a	627 of	er of			SEQ ID	Je .		AW024164, C06355, AI476066,	nore C06056, R78935, AI436456, AI064830, AL121270	AL047042, AL046849, AI349772, AI686926,	AL045500, AI433157, AL047763,	
SEQ ID NO:1391, b is an integer 15 to 423, where both a and b	collespond to the positions of nucleotide residues shown in SEQ	NO:1391, and where b is greater	than or equal to a + 14.	Ω	present invention are one or more	polynucleotides comprising a	nucleotide sequence described by	the general formula of a-b, where	is any integer between 1 to 84	SEQ ID NO:1392, b is an integer	15 to 856, where both a and b	correspond to the positions of	nucleotide residues shown in SEQ	NO:1392, and where b is greater	than or equal to a + 14.	Preferably excluded from the	present invention are one or m	polynucleotides comprising a	nucleotide sequence described by-	of a-b,	is any integer between 1 to 62	SEQ ID NO:1393, b is an integer	15 to 641, where both a and b	correspond to the positions of	nucleotide residues shown in SEQ	NO:1393, and where b is greater	than or equal to a + 14.	Preferably excluded from the	present invention are one or more	polynucleotides comprising a	nucleotide sequence described by	
				876201												876206		•					•					876207				
				HISAQ01												HCRMC10												HWABD53				_
				1392												1393												1394				

ALO36980, ALO36146, d to the positions of e residues shown in SEQ ID AI34937, AI538716, and where b is greater AW089572, AI497733, AI34064, AI597750, AI863014, AI39771, AI440239, AI800433, AW238730, AI80433, AW238730, AI80433, AW238730, AI866608, AI281773, AI69924, AI697137, AI909641, AI697137, AI69719, AI697137, AI69719, AI697137, AI69719, AI697137, AI69719, AI686608, AI286608, AI286608, AI386608, AI686608, AI686608, AI686608, AI686608, AI686608, AI686608, AI697137, AI69		SEO ID NO:1394. b is an integer of	AL119791,	A1440426,	AIS00077,	AI281779,	
d to the positions of d to the positions of e residues shown in SEQ ID AI349937, AI538716, AI349645, AI349937, AI538716, AI340682, AM071417, AI34904, AI597750, AI34064, AI597750, AI34904, AI597750, AI34036, AW1925750, AI34036, AW19192, AI34036, AW193152, AI34036, AW193162, AI34036, AW193087, AI56870, AW274192, AI637137, AI440239, AI637137, AI440239, AI637137, AI39087, AI637137, AI609641, AI637137, AI609641, AI697137, AI699641, AI69660, AI697176, AI69660, AI69660, AI69660, AI69660, AI69660, AI697176, AI69660, AI69660, AI69660, AI69660, AI6971776, AI69660, AI6960, AI6	_	, where both a	98	AL036146,	AW074993,	A1687728,	
e residues shown in SEQ ID A1349937, AL19748, and where b is greater A1340064, A1538716, A1340064, A1597750, A1349004, A1597750, A1349004, A159750, A1863014, A1590128, A1521012, A1282655, A1036396, AM195957, A1036396, AM195957, A1036396, AM195957, A1036396, AM195957, A1036396, A1440239, A1036396, A14439087, A1036396, A1639087, A1036396, A1699137, A10314902, A1699137, A10314902, A1699137, A10314902, A1699257, A1036328, A1699257, A1036328, A1692540, A1889203, AW301300, A1889203, AW301300, A1366549, A1612913, AA572758, AL036759,		d to the positions	883	4	AW268253,	AI312152,	
and where b is greater Augual to a + 14. Augual t		e residues shown in SEQ	573	4	AIS67351,	AI620284,	
AW089572, AI497733, AI340582, AW071417, AI349004, AI597750, AI863014, AI590128, AI521012, AI282655, AL036396, AW195957, AI568870, AW274192, AI702406, AW303152, AI702406, AW303152, AI702406, AW303152, AI702406, AW303152, AI800433, AW238730, AI834036, AI679724, AI631107, AI631107, AI697137, AI631107, AI697137, AI631107, AI6906328, AI669592, AI702433, AI687375, AI702433, AI687375, AI899203, AW301300, AI534902, AI283941, AA585422, AI492540, AI889203, AW301300, AI534907, AI366549, AI612913, AA572758, AL036759,		and where b is	993	AI538716,	AI469532,	AI699857,	
AI340582, AW071417, AI349004, AI597750, AI863014, AI590128, AI521012, AI282655, AL036396, AW195957, AI568870, AW274192, AI702406, AW303152, AW103371, AI440239, AI800433, AW238730, AI934036, AI639087, AI635461, AI439087, AL121365, AI635942, AL281773, AI9909641, AI34902, AI445432, AI5906328, AI69592, AI500659, AI445432, AI500659, AI69557, AI702433, AI687375, AI702433, AI687375, AI889203, AW301300, AI539771, AW167776, AI5664079, AA613907, AI366549, AI612913, AA572758, AL036759,		equal to a + 14.	957	9773	AI818683,	AW169653,	
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9, AI612913, 8, AL036759,				σ	AI909666,	AI673256,	
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AI340519, AI690751, AI			I34051	69075	AI349226,	AI568854,	

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			AC006039, AL137429, AF026124,
			I42402, AR013797, AC007392, AL035067, AC007172,
			AL133077, I26207, AL137526, AL137560, E15569,
			AC004200, AJ012755, AL050172, AC004690,
			AF100931, A93350, I66342, AL137533, AL035587,
			, U39656, AF026816,
			AC005291, AC004383, I00734, AF057300
1395 HKCSF17	17 876208	Preferably excluded from the	
		present invention are one or more	
		polynucleotides comprising a	
		nucleotide sequence described by	
		al formula of a-b, wher	
		teger between 1 to 906	
		SEQ ID NO:1395, b is an integer of	

	15 to 9 corresp nucleot NO:1395 than or present polynuc nucleot the gen is any SEQ ID 15 to 1 corresp nucleot the gen is any SEQ ID 15 to 4 corresp nucleot the gen is any SEQ ID 15 to 4 corresp nucleot the gen is any SEQ ID prefera present polynuc nucleot the gen is any SEQ ID 15 to 4 corresp nucleot the gen is any SEQ ID 15 to 4 corresp nucleot the gen is any SEQ ID 15 to 4 corresp	15 to 920, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1395, and where b is greater than or equal to a + 14.	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 1087 of SEQ ID NO:1396, b is an integer of 15 to 1101, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1396, and where b is greater than or equal to a + 14.	Preferably excluded from the present invention are one or more present invention are one or more present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 434 of SEQ ID NO:1397, b is an integer of 15 to 448, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1397, and where b is greater than or equal to a + 14.	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a AI088609, AI742316, AI229474, AA442089, AI886144, AI307145, AI129474, AA442089, AI886144, AI307145, AI307146, AI307
876213			HTDA112	HYABB57	HWLVN09
	HTDAI12 HYABB57 HWLVN09		1396	1397	1398

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			15 to 762 whore both a and h		
			correspond to the positions of		
			-		
			than or equal to a + 14.	- 1	
1399	HOHAU02	876220	Preferably excluded from the		420, AC005082,
			present invention are one or more	AC008064, AL022727	
	-		polynucleotides comprising a		
			nucleotide sequence described by		
			the general formula of a-b, where a		
			is any integer between 1 to 305 of		
			15 to 319, where both a and b		
			correspond to the positions of		
			nucleotide residues shown in SEQ ID		
			NO:1399, and where b is greater		
			equal to a + 14.		
1400	HCRNJ43	876224	Preferably excluded from the	AA313797, W73983, AW374097	7, AA824282, AI207345,
			present invention are one or more	226317	
			_		
			nucleotide sequence described by		
			15 to 1575, where both a and b		
			correspond to the positions of		
			nucleotide residues shown in SEQ ID		
			NO:1400, and where b is greater		
			than or equal to a + 14.		
1401	HWLGV14	876226	Preferably excluded from the	AI110653, AA573785, AI42182	829, AI889106,
			present invention are one or more	AI815098, AW082282, AW151910	.910, AA309046,
			polynucleotides comprising a	AW251068, AI688082, AI935867,	867, AA903732,
			nucleotide sequence described by	AI342309, AI469758, AI301940	.940, AI336447,
			the general formula of a-b, where a	AI660665, AI625318, AI636809,	809, AI559518,
			is any integer between 1 to 1299 of	AI216199, AA974182, AI336445	445, AI476296,
			SEQ ID NO:1401, b is an integer of	AI272699, AA865622, R95048	8, AI832439, AI908555,

			15 to 1313, where both a and b	AW079674, AW276067, H71284, AI290972, AI659188,
			correspond to the positions of	H41084, H39231, AI865986, AI333305, R76336,
_			nucleotide residues shown in SEQ ID	AI914585, AI590410, H12385, AA987621, R48364,
			NO:1401, and where b is greater	R94963, AA639087, D45438, C20912, AI274107,
			than or equal to a + 14.	AI720940, H70884, AA372940, AW250334, H15022,
				AI244423, AW192993, AA935031, AI199655,
				, H15021, A
				AA886276, AI2255252, R45920, AF115384, AC006479
1402	HCYBM15	876228	Preferably excluded from the	AA305646, D57483, C14389, D80391, D59787,
			present invention are one or more	D80196, D81026, D80253, D80522, D58283, D80366,
	-		polynucleotides comprising a	D51022, D80227, D59859, D59467, D80043, D51423,
	2		nucleotide sequence described by	D80022, C14331, D59275, D80166, D80195, D59619,
			the general formula of a-b, where a	D80210, D51799, D80164, D80240, D59927, D59502,
	-		is any integer between 1 to 516 of	D81030, D50979, D59889, D80248, D80212, D80251,
			SEQ ID NO:1402, b is an integer of	D50995, D80269, D80188, D80219, C15076, D80038,
			15 to 530, where both a and b	AA305578, D80133, D59610, D80024, AA305409,
	•		correspond to the positions of	AA514186
_			nucleotide residues shown in SEQ ID	C75259,
			NO:1402, and where b is greater	
			than or equal to a + 14.	D80132, C14014, D58253, AW378532, AW375405,
	•			
_				
				AW369651, AW177505, AW179024, AW352158, F13647,
_				3
_				AW377676, D80247, AW178906, AW352170, AW177731,
				AW179019, AI910186,
				AW178980, AW177733, AW378528, AW178908,
				AW178754, D51079, AW179018, AW352174, AW179004,
				AW378525, C06015, AW367967, D80157, AW177722,
				D51759, AW178774, AW178911, AW378543, AW352163,
				AW378540, AW178983, Z21582, AW178781, T48593,

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D59653, C14227, D51213, D45260, AW352120,
C14227, D51213, D45260,
71177 TE1213 DAE260

			15 to 1410 those both 1 12 th	
			nd to the positions o	
			s shown ir	
			NO:1403, and where b is greater	
			than or equal to a + 14.	
1404	HHFCN93	876232	0	AA769099, AW051928, AI701149, AW166012, H14423,
			present invention are one or more	AA972142, AI339332, N92764, R59745, AA100558,
			polynucleotides comprising a	AI383947, AA347767, AA015757, AI338203,
			nucleotide sequence described by	AA347768, D81417, H72916, AA805417, D20390,
			the general formula of a-b, where a	
			is any integer between 1 to 1428 of	
			SEQ ID NO:1404, b is an integer of	
			15 to 1442, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1404, and where b is greater	
1405	H2CBC05	876236	Preferably excluded from the	AI743549, AI953907, AW444710, AI457576,
			present invention are one or more	AA452352, AI744355, AW169608, AA452129,
			polynucleotides comprising a	AA809771, AI284062, AA307160, AW363101,
			nucleotide sequence described by	AI865348, AA907553, AI620087, AI936509,
			the general formula of a-b, where a	AA618311, AA456277, AA454662, AA173381,
			is any integer between 1 to 1675 of	
			SEQ ID NO:1405, b is an integer of	AW363100, AA478933, N90372, AI186424, C14331,
			15 to 1689, where both a and b	D80166, AA809122, D80439, D80247, D59619,
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	D80022, D81030, D80219, D80212, D80133, C14389,
			NO:1405, and where b is greater	, D59787
			than or equal to a + 14.	6, D59502, D51423, D51799,
				DS1060,
				D80248,
				D80188, D80164, D59467, D51022, D59275, D80038,
				D50995,
				D80269, D59889, D59653, D50979, D80024,
				78, D51759, D80302, AA514188
				T03269, D80241, D80251, AI535686, AW377671,

C14407, C AW17501, AW179328, AW179328, AW178705, AW1788, YY1188, YAN188, YAN18	AIS25923, C05695, AW178893, D45260, C75259, D58246, D59373, AW375405, AW360844, H67866, C14407, C03092, H67854, C14973, AW366296, AW177501, AW179328, AW177511, AW360817, AW179328, AW179020, T48593, AW375406, AW378534, AW179328, AW179020, T48593, AW37672, AW179023, AW178905, D80064, AW177731, AW378528, AW178762, AW178754, AW179019, AW177731, AW378528, AW177505, AW178754, AW179019, AW177731, AW37004, D59503, AW177731, C14227, AW177505, AW178907, AW178907, AW178908, AW1779018, AW352158, AW178909, AW17734, AW378533, D80949, AA514184, AW352170, D592117, D59317, D80014, D59474, N66429, AI525920, AW177734, AW178009, AW178911, AW378533, D80949, AA514184, AW360834, AW378533, D80949, AA514184, AW378530, D58101, AW178774, AW178781, AW378543, AW378543, AW378539, AI525212, AW378525, C144967, D60010, H67858, AW178531, D59055, D59657, D51033, D45273, AM178781, AW378539, AI525222, T02974, D13645, A62298, AR6916, A86595, AR565222, T02974, D13645, A62298, AR69166, A86596, AR060385, AR0828138, A62300, A30438, AR008277, AR008281, Y17187, AR008281, AW17818, Y17187, AR008278, AR060385, AB028859, AJ132110, AB002449, I50126, I50132, I50128,
A45456,	A94995, D26022, A26615, ARO
A43192,	Y12724, A43190, AR038669, A
AR0664	88, Y09669
D89785	, A78862,

A63261, D88547, D50010, AR062872, A70867, AR008408, A64136, A68321, I79511, AR025207, D13509, AR060133, AF123263	rably excluded from the high energy in the formula of a-b, where a high energy integer both a and b spond to the positions of the confidence of the high energian in SEQ ID off, and where b is greater or equal to a + 14.	rably excluded from the nt invention are one or more AB010812, AC004520, AF125534, AC007225 acclerate sequence described by eneral formula of a-b, where a y integer between 1 to 824 of D NO:1407, b is an integer of 838, where both a and b spond to the positions of otide residues shown in SEQ ID otide residues at 14.	rably excluded from the AI017564, AA809290, AW002023, AA405338, at invention are one or more AA806993, AA405339, AA888974, AA236935, acleotides comprising a AI024655, AA262702, H49789, AI524770, N77703, aneral formula of a-b, where a y integer between 1 to 918 of 932, where both a and b shood to the positions of
	m the on ing a cribe cribe cribe cribe a-b, a-b, inte and ions wn ir.	rom the ising a lescribed a-b, wen 1 to 8 an integ an integina of those is great is great 14.	Preferably excluded from the present invention are one or polynucleotides comprising a nucleotide sequence described the general formula of a-b, wlis any integer between 1 to 93 SEQ ID NO:1408, b is an integer to 932, where both a and by the positions of the po
	876238	876239	876259
	HTEPE28	HUSGL79	HPMFU84
	1406	1407	1408

			The standard residing shown in SEO ID	
			NO.1400 and thorn but and the telestation	
			NO:1400, and where D is greater than or equal to a + 14.	
1409	HDLAD09	876260	Preferably excluded from the	W79877, Z42158
			present invention are one or more	
			polynucleotides comprising a	
			nucleotide sequence described by	
			is any integer between 1 to 751 of	
			SEQ ID NO:1409, b is an integer of	
			15 to 765, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1409, and where b is greater	
			than or equal to a + 14.	
1410	HCQAW45	876261	Preferably excluded from the	AI829532, AL008582
			present invention are one or more	
			polynucleotides comprising a	
			nucleotide sequence described by	
			the general formula of a-b, where a	
			is any integer between 1 to 518 of	
			SEQ ID NO:1410, b is an integer of	
			15 to 532, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1410, and where b is greater	
			than or equal to a + 14.	
1411	HCYAC01	876265	Preferably excluded from the	14, AA308913, D59927, D50979, D802
			present invention are one or more	D58283, D80188, D80253, D80195, D80043, D59275,
			polynucleotides comprising a	D80269, D59502, D59859, D80022, D80166, D80366,
			nucleotide sequence described by	D81030, D51423, D59619, D80210, D51799, D80391,
			the general formula of a-b, where a	D80240, D59787, D80378, D80038, D80212, D80045,
			is any integer between 1 to 538 of	D80193, D80196, D80164, D80219, D57483, C14389,
			SEQ ID NO:1411, b is an integer of	D59889, D50995, D80024, D59467, D59610, C14331,
			15 to 552, where both a and b	C15076, C14429, D80241, D51060, AA305409,
			correspond to the positions of	T03269, D80522, D58253, C75259, C14014,

		nucleotide residues shown in SEQ ID	AW178893, D81026, D80134, AA305578, D51022,
		, and where b	AW179328, D51250, D80268, AW177440, F13647,
		than or equal to a + 14.	', AW178775, D80251, D80949
	_		D59695, AA514188, D52291,
,-			352158, D80248,
			AW178762, AI905856, AW177501, AW177511,
			D80064, C05695, AW352117, C14407, AW176467,
			AW375405, AW377671, D80132, AW360834, AW378540,
			AA285331, AW366296,
			AW375406, AW378534, D51097, AW179332, AW377672,
			AW179023, AW178905, AW352171, AW377676, D80439,
	,		AW178906, AW352170, AW17731, AW178907,
			AW179019, AW179024, D59373, D80247, AW179220,
			AW177505, AW360841,
	_		
			AW178908, AW178754
_			D51103, AW
			AW352174, AW178914, T03116, AW378525, AW367967,
	-		W177722, D
			AW178774, AW178911, AW378543, AW352163, D80258,
			, D59653
			l, D45260, T4
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			, AA514184, AI525917, AI5252
			_
			, AI535686, D51221,
			D58101, AI
			C16955,
			AW378542, C05763, Z33452, AI525522, AW360855,
			, D31458, CO
			13958, U49017, A84916, AJ132110,
			298,
			A25909, A67220, D89785, A78862, D34614, D88547,

				ARONAS78 AFOSR696 X82626 ARO28859 182448
				, Y12724, AB012117, A82595, X6
				AB002449, AR060385, AR016808, A85396, AR066482,
				A44171, A94995, A85477, I19525, A86792, U87250,
				AR008443,
				I50133, AR016514, AR066488, AR060138, A45456,
				A26615, AR052274, I14842, Y09669, AR066487,
				A43192, A43190, AR038669, AR054175, A30438,
				AR066490, Y17187, I18367, A63261, AF135125,
				D88507, AR008277, AR008281, D50010, AR062872,
_				A70867, AR016691, AR016690, U46128, AR008408,
				I79511, A64136, A68321, AB033111, D13509,
				U87247, AR060133, AR064240, AF123263, AR032065,
				U79457, X93535
1412	HCROF86	876266	Preferably excluded from the	AI650543, W69438, W69521, H10084, AA489949,
			present invention are one or more	13027, F079
			polynucleotides comprising a	AW388196, AW388234, AW388225, AW388262,
			nucleotide sequence described by	AW388176, AW388206, AW388208, AW388214,
			the general formula of a-b, where a	AW388253, AF086275, AB024057, AB017114, U88873
			SEQ ID NO:1412, b is an integer of	
			15 to 1100, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1412, and where b is greater	
			than or equal to a + 14.	
1413	H2CBJ83	876269	Preferably excluded from the	AA403070, AA313305, AA361460, T78498
			present invention are one or more	
			polynucleotides comprising a	
			nucleotide sequence described by	
			al formula of a-b, wher	
		_	is any integer between 1 to 549 of	
			SEQ ID NO:1413, b is an integer of	
			15 to 563, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	

				8, X67155, Y17188, D26022, A25909
				A67220, D89785,
				AR060385, AR025207, AB002449, AR008443, I50126,
				I50132, I50128, I50133, AR066488, AR016514,
				AR060138, A45456, A26615, AR052274, I14842,
				Y09669, A43192, A43190, AR038669, AR054175,
				AR066487, AR062872, A30438, Y17187, X68127,
				A63261, D50010, AR008277, AR008281, A70867,
				9
				A64136, A68321, AR060133
1415	HWMCL22	876274	Preferably excluded from the	R86344, R86183, AC004686
			present invention are one or more	
			polynucleotides comprising a	
			nucleotide sequence described by	
			the general formula of a-b, where a	
			15 to 418, where both a and b	
			correspond to the positions of	
			residues	
			NO:1415, and where b is greater	
	:		than or equal to a + 14.	
1416	HCRPZ42	876276	Preferably excluded from the	AA285061
			present invention are one or more	
			polynucleotides comprising a	
			nucleotide sequence described by	
			al formula of a-b, wher	
			is any integer between 1 to 499 of	
			SEQ ID NO:1416, b is an integer of	
			15 to 513, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1416, and where b is greater	
			than or equal to a + 14.	
1417	HCYBM32	876277	Preferably excluded from the	AA305407, D51423, D51799, D80166, C14389,

	present invention are one or more	D80522, D81030,
	polynucleotides comprising a	D80366, D81026, D59859, D59619, D80210, D80240,
	nucleotide sequence described by	AW377671, D80269, C14331, D582
	the general formula of a-b, where a	D80212, D50995, D80188, D59467, D51022, D80022,
	is any integer between 1 to 428 of	AA305409,
	inte	1, D80164, D59275,
	15 to 442, where both a and b	D59502, D80241, D80251, D57483, D59889, D80196,
	d to the po	4, D59927, AA514188
	$\boldsymbol{\sigma}$	8, D80193
	NO:1417, and where b is greater	D80378, D80439, AW360811, AW177440, C14429,
	than or equal to a + 14.	i, C75259
		D80247, T03269, D80302, AW375405, AW360844,
		3
		AW360817, AW375406, AW178906, AW378534,
	-	AW352171, AW179332, AW377672, AW179023, D80157,
		AW178905, C05695, AW378532, AW377676, D51103,
		AW360834, D51759, D80134, AW177505, AW360841,
		AW352170
		, AW178907,
		AW179018,
		AW176467, AW369651, D45260, AW179020, AW177456,
		1179329, AW
		AW178908, AW17897;
		AW179004, AW178774,
		AW179009, AW179012, D80064, D80258, C14227,
-		AW352120, AW378525,
		H67854, C14077, D50981, D58246, C03092,
		AI525923, T02974, AW178911, H67866, AW17722,
_		AI910186, AW17728, AA514184, AA809122, T03116,
		D59503, AW367950, AI905856, AW378540, D59317,
		AI525917,
		D45273, D51221, T03048, D60214, C14344, D59474,
		AW178986, C14973, AW378533, AIS57774, AI535850,
		39, AW177734, AW177723, C14957
		C14298, AI535686, AI525235, D59551, AI525215,

112, AW179013, D51213, 125, Z21582, D51097, 10160, AW378542, AI525928, AW360855, D51053, C04682, C015606, D50313, E12830, A62298, A84916, A62300, S07, A82595, AR008443, D26022, Y12724, A67220, D89785, C0132, I50128, A66487, I14842, AR066480, AR08277, I18367, X68127, I18367, X68127, I18367, X68127, AR016690, U46128, A68321, I79511, AR0166482, A85396, AR032065, A85477, AR032065, A85477, A8132065, A81382	A1910457, AA496921, AW082592, AI650301, 010258, U81511,
AM179011, AI525912, AA285331, AI525925, C16955, Z33452, Z30160 D80949, AW178759, AI520, D59695, D52291, D510 T02868, D50312, AF01566, D50314, D88159, E1286, AB0028859, AF015607, X67155, Y17188, D260 AB002449, A94995, A672 AB002449, A94995, A672 AB002449, A94995, A672 A343190, AR016691, AR066, A30438, Y17187, AR066, A30438, Y17187, AR066, A30438, Y17187, AR066, A1190, AR016691, AR111, A113263, AR060133, AF123263, A4111, AR03 A86792, U79457, X93545	AI246769, AI304342, AA541292, AI129972, AA627519, AA627188, AA577580, AW439990, AC004080, U41813, AF 13536, M28449
AI52522, AI52522, AI52522, H67858, C1 C05763, D8 AI525237, C06084, TC AF018138, AJ132110, AR018138, AJ132110, AR060385, A25909, AE A78862, D3 D88547, AF A26615, AA AR062811, AR062872, AR008281, AR062872, AR008281, AR012117, D88507, AI	AI346422, AI381007, AW089855, AM923632, AI676154, XI3537, X
·	preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 915 of SEQ ID NO:1418, b is an integer of 15 to 929, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1418, and where b is greater than or equal to a + 14.
	876278
	HCRPJ72
	1418

			present invention are one or more		
			leotides comprising a		
			nucleotide sequence described by		
			the general formula of a-b, where a		
			is any integer between 1 to 230 of		
			SEQ ID NO:1419, b is an integer of		
			15 to 244, where both a and b		
			correspond to the positions of		
			nucleotide residues shown in SEQ ID		
			NO:1419, and where b is greater		
			than or equal to a + 14.		
1420	HMWFC49	876281	Preferably excluded from the	AW410053	
			present invention are one or more		
			polynucleotides comprising a		
			nucleotide sequence described by		
			the general formula of a-b, where a		
			is any integer between 1 to 158 of		
			SEQ ID NO:1420, b is an integer of		
			15 to 172, where both a and b		
			correspond to the positions of		
			nucleotide residues shown in SEQ ID		
			NO:1420, and where b is greater		
			than or equal to a + 14.		
1421	HMSIE02	876282	ρ	AW451452, AI040326, AI650832	832, AA313243,
		•	present invention are one or more	AI650393, AI818259, AA534633	633, AI094737,
			polynucleotides comprising a	AI033652, AI693411, AI341518	518, W30723, AW197245,
			nucleotide sequence described by	AW051598, AW291994, AI274289	_
			the general formula of a-b, where a	AA035621, AA653321, AA634950	_`
			is any integer between 1 to 2279 of	AA136077, N99062, AA806117,	7, AA136161, AA722867,
			SEQ ID NO:1421, b is an integer of	AA932876, AI435016, AI659053	053, AI474321, H87560,
			15 to 2293, where both a and b	AA843369, H21542, AA361623,	N47604, N4
			correspond to the positions of	AI907694, AA332538, H87452,	
			nucleotide residues shown in SEQ ID	AA365059	
			, and		
			than or equal to a + 14.		
1422	HCRMZ34	876284	Preferably excluded from the	AA034416, AA491400, AA504783	783, W65331, AI885434,

			present invention are one or more polynucleotides comprising a	AI553873, AI637992, AW172551, AA236838, AA053881, AA482166, AI680567, AI184074, R43006
			nucleotide sequence described by	
			the general formula of a-b, where a	096876
			Ψ	
			SEQ ID NO:1422, b is an integer of	
			15 to 1660, where both a and b	
			d to the positions of	
		_	nucleotide residues shown in SEQ ID	
			NO:1422, and where b is greater	
			than or equal to a + 14.	
1423	HTGAM27	876300	Preferably excluded from the	AA187449, AW361774, AL034396, L14787, Z99130,
			present invention are one or more	AL031115
			polynucleotides comprising a	
			nucleotide sequence described by	
			the general formula of a-b, where a	
			is any integer between 1 to 296 of	
			SEQ ID NO:1423, b is an integer of	
			15 to 310, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1423, and where b is greater	
			than or equal to a + 14.	
1424	HCYBI20	876304	Preferably excluded from the	, AI763355,
			present invention are one or more	AI609777, AI859398, AA197062, AA305389,
			polynucleotides comprising a	, AW271204, AA825907,
			nucleotide sequence described by	AI910841, AI673503, AI632367, AW269183,
			the general formula of a-b, where a	AW196356, AW273255, AI304550, AI419935,
			is any integer between 1 to 3092 of	AI270299, AI247514, W01219, AI355117, N72988,
			SEQ ID NO:1424, b is an integer of	AA030042, AW007158, AA070475, AW006961,
			15 to 3106, where both a and b	AI304462, W57671, AA876039, AA705874, AA831500,
			correspond to the positions of	H62242, AA897761, W03289, AA029912, AA305307,
			nucleotide residues shown in SEQ ID	H93491, W91963, H82187, AI245415, AA643520,
			NO:1424, and where b is greater	AW088307, H93492, R89908, AA377111, AI318375,
			than or equal to a + 14.	AI961885, AA059231, AA883186, AW139085,
				AA581261, T85676, Z40302, AA887782, AA502293,

				, H62331, R93209, R07861,
				H82082, T29678, F01458, AA527320, H61166,
				AI270229, AI932770, AW070350, R07916, AI765901,
				F04303, N74218, AA581216, AW268185, AI334444,
				AW274341, AW268947, AA128235, AI699588,
				851,
				œ
				D59502,
				, D80166, D80195, D51423, D5961
				D80391, D80164, D59275, D80240, D80253, D80038,
				O
				D81026, D80212,
				D80196, D80188, D51022, D50979, D80219, D80378,
				AA305578, AA5141
	_			C03092, D59889, D80193, D80133, D80045,
				6, D80024
				D80157, AW360811, D51103, AW177440, D59653,
				D51759, D80241, D80251, AW178893, T03269,
				71, AW375405, C75259, H67866
				AW360844
				A26237,
				0, AR0181
	_			7
				5, AR060138
		•		, D26022,
				Y12724, A43190, AR038669, A25909, AR066488,
				Y09669, AR066487, Y17187, A67220, D89785,
				A78862, D34614, A30438, AR008443, A63261,
				AR008277, AR008281, AR062872, A70867, AR016691,
				AR016690, U46128, D50010, D88547, I79511,
				64136, A68321, AR008408,
				AR025207, AR060133, AF123263, AR032065
1425	HNEDH18	876306	Preferably excluded from the	AA504982, AL11940
			present invention are one or more	M20317, X1

1426 HWMFQ61	Q61 876308	the general formults any integer bet SEQ ID NO:1425, b 15 to 352, where be correspond to the nucleotide residue NO:1425, and where than or equal to a Preferably exclude present invention polynucleotides conucleotide sequenct the general formults any integer bet SEQ ID NO:1426, b	2, AA524145, AW007155, 6, AI815931, AW193517, 8, AA573859, AI879177, 6, AI376231, AI352472,
		the general formula of a-b, where is any integer between 1 to 338 o SEQ ID NO:1425, b is an integer of 15 to 352, where both a and b correspond to the positions of nucleotide residues shown in SEQ NO:1425, and where b is greater than or equal to a + 14. Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where is any integer between 1 to 1953 SEQ ID NO:1426, b is an integer o	2, AA524145, AW007155, 6, AI815931, AW193517, 8, AA573859, AI879177, 6, AI376231, AI352472,
		is any integer between 1 to 338 o SEQ ID NO:1425, b is an integer o 15 to 352, where both a and b correspond to the positions of nucleotide residues shown in SEQ NO:1425, and where b is greater than or equal to a + 14. Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where is any integer between 1 to 1953 SEQ ID NO:1426, b is an integer o	2, AA524145, AW007155, 6, AI815931, AW193517, 8, AA573859, AI879177, 6, AI376231, AI352472,
		SEQ ID NO:1425, b is an integer of 15 to 352, where both a and b correspond to the positions of nucleotide residues shown in SEQ NO:1425, and where b is greater than or equal to a + 14. Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where is any integer between 1 to 1953 SEQ ID NO:1426, b is an integer of a sequence of a sequen	2, AA524145, AW007155, 6, AI815931, AW193517, 8, AA573859, AI879177, 6, AI376231, AI352472,
		15 to 352, where both a and b correspond to the positions of nucleotide residues shown in SEQ NO:1425, and where b is greater than or equal to a + 14. Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where is any integer between 1 to 1953 SEQ ID NO:1426, b is an integer o	2, AA524145, AW007155, 6, AI815931, AW193517, 8, AA573859, AI879177, 6, AI376231, AI352472,
		correspond to the positions of nucleotide residues shown in SEQ NO:1425, and where b is greater than or equal to a + 14. Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where is any integer between 1 to 1953 SEQ ID NO:1426, b is an integer or	2, AA524145, AW007155, 6, AI815931, AW193517, 8, AA573859, AI879177, 6, AI376231, AI352472,
		nucleotide residues shown in SEQ NO:1425, and where b is greater than or equal to a + 14. Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where is any integer between 1 to 1953 SEQ ID NO:1426, b is an integer or sequence of the	2, AA524145, AW007155, 6, AI815931, AW193517, 8, AA573859, AI879177, 6, AI376231, AI352472,
		NO:1425, than or e Preferabl present i polynucle nucleotid the gener is any in SEQ ID NO	2, AA524145, AW007155, 6, AI815931, AW193517, 8, AA573859, AI879177, 6, AI376231, AI352472,
		than or e Preferabl present i polynucle nucleotid the gener is any in SEQ ID NO	2, AA524145, AW007155, 6, AI815931, AW193517, 8, AA573859, AI879177, 6, AI376231, AI352472,
		Preferabl present i polynucle nucleotid the gener is any in	2, AA524145, AW007155, 6, AI815931, AW193517, 8, AA573859, AI879177, 6, AI376231, AI352472,
	,	present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 1953 of SEQ ID NO:1426, b is an integer of	6, AI815931, AW193517, 8, AA573859, AI879177, 6, AI376231, AI352472,
		polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 1953 of SEQ ID NO:1426, b is an integer of	, AA573859, AI879177, AI376231, AI352472,
		nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 1953 of SEQ ID NO:1426, b is an integer of	, AI376231, AI352472,
		the general formula of a-b, where a is any integer between 1 to 1953 of SEQ ID NO:1426, b is an integer of	CONTRA COURT
		is any integer between 1 to 1953 of SEQ ID NO:1426, b is an integer of	A19561/2, AA283/02, AA5834/9, AA486429,
		SEQ ID NO:1426, b is an integer of	AI095623, N91996, AA405889, AI089975, AA493377,
			AI147623, AA147930, H09366, AI879560, AI698813,
		15 to 1967, where both a and b	AI493913, AA580211, AA737974, AI476337,
	_	correspond to the positions of	AA423896, N24051, N32340, N66204, AA405729,
		nucleotide residues shown in SEQ ID	AA507484, AI374680, AA489431, AA157554,
		NO:1426, and where b is greater	AA147501, N35409, AA505515, AA489372, AA127433,
		equal to a + 14.	N55519, H15112, AA173145, N57433, AA471177,
			AW401453, N63852, T78215, AA857801, N52066,
			H09309, AA780883, AL079771, AA356048, AA769879,
			AA173273, R25268, AA127432, R46621, AI707462,
			AA807765, AI423315, AA877529, AA836375,
			AA352973, AA148410, H85254, AA356047, AA326793,
			AA678778, R53945, AA278977, N99204, AA335034,
			R07396, AA423831, AA367574, AA715745, H84922,
			AI762734, R07347, F05138, AA058460, AW339712,
			AI701737, T29480, AA995682, AI815735, N48041,
			AI362375, N35874, F01382, AA329166, AA295203,
			, AW020406,
			AL041150, AW020397, AI491904, AI564716,

AI923989, AW021717, AW410302, AI224373,	AI307557, AA464646, AW020592, AI289310,	AI859991, AW236692, AI60976	AI879064, AI267185, AI567582, AL042753,	AW020095, AI811603, AI621341, AI311472,	AL038986, AI049850, AI927233, AI656188,	AI560722, AA806534, AA502794, AI350489,	679506, AW020710, AI961	AI580214, AL048871, AI349012, AI521005,	AL079963, AL036705, AI525653, AI581033,	AIS90943, AI758445, AA580663, AI432570,	ω,	AI242248, AI741158, AI499963, AW102798,	,	AI638644, AI537677, AI434731, AW148478,	AI141727, AW020373, AL048323, AI432507,	AW169784, AL048340, AI382313, AI587209, N22276,	AA514684, AI282268, N29277, AI538764, AI440263,	AW020419, AI587000, AW160905, AW162194,	,	, AI499279,	, AI697236,	AI348901, H41759, AI500061, AI372009, AW327825,	8, AA455772, AI699865,	, AI279925,	6, AL046466,	6, AI471282, AI500514,	, AI950892, AI341690, AW05108	, AI624245, AI5246	AI472484, AW265582, AI698391, AI538564,	AL036361, X15653, Y09008, A64377, AC007637,	X89398, AC010582, Y08975, X99018, U55041,	X92986,	7, I89947,	, AL1220	AL137533. A08910. A08909. AL117460 AF026124
										-				-																					_

	AF145233, A08908, Y11254, AL133560, AF082526,	
	M85164, X70514, AL049996, AL050172, AJ005690,	
	AR038854, AL110296, AF090900, AL080156,	
	U91329, J05277, AL049283, AF087943, A08912,	
	AF146568, AF113690, AL133080, U42766, S76508,	
	E06788, E06790, E06789, AF061795, AF151685,	
-		
)5822, AL133640, AL049347	
	AF118094, X06146, Y09972, E12747, A21103,	
	X99257,	
	AL137530, A76335, AR038969, AF111851, X63162,	
	AF079763, AF111849, AL137574, S77771, S83440,	
	S68736, A08911, AL080118, A18777, AL122110,	1
	AF061943, X67688, Y16645, AL110218, AF113699,	
	AF069506, AF141289, U86379, I48979, AJ010277,	
	I89931, A77033, A77035, AF017790, Z72491,	
	A08907, L04849, AF065135, AF081366, S69385,	_
	AL133016, AJ003118, AL096728, AL050280, U55017,	
	9, AL110269, A15345, AL1176	
	AF067728,	_
	.122093, Y07905,	
		-
	AL049339,	
	AF106657	_
	U02475, Y10936, AL	
-	M27260, AL023657,	
	E02319, I3	
	AL137479, AC002467, AL122049	
	AL117416, U95114, X92070, AL137254, AL080074,	

				- 1
				AL049452, I32738, A23630, AF077051, AL110159,
				X63410, Y10655, S63521, AL049300, A86558,
				AF090943, X79812, AL110196, AF176651, X84990,
				AB007812, E01314, Z37987, AL133075, A07647,
				AF124728, AF036268, AL122045, I66342, AL050146,
				, AL133619,
				AR053103, AC004878,
1427	HFIUZ10	876309	Preferably excluded from the	AA449704,
			present invention are one or more	AA448557, AI453006, AA863038, AI277552,
			polynucleotides comprising a	AA723892, AI282002, AA879085, AI282089,
			nucleotide sequence described by	AA928469, T81791, AA258329, AI271667, R02362,
			the general formula of a-b, where a	T82108, H66854, AC004080, M74297
			is any integer between 1 to 865 of	
			SEQ ID NO:1427, b is an integer of	
			d to the positi	
			nucleotide residues shown in SEO ID	
			and where b is greater	
			equal to a + 14.	
1428	HDPJE43	876322		AA305011, M73047, X81323, U50194, A58393.
				A58395
			leotides comprising a	
			nucleotide sequence described by	
			al formula	
			teger between	
			SEQ ID NO:1428, b is an integer of	
			15 to 521, where both a and b	
		_	correspond to the positions of	
			nucleotide residues shown in SEQ ID	
		_	NO:1428, and where b is greater	
			than or equal to a + 14.	
1429	HWLWR2	876326	Preferably excluded from the	AW291224, AA027791, AI826645, AI970074, AI859242
	2		present invention are one or more	
			polynucleotides comprising a	
			nucleotide sequence described by	

1432 HWLQG81	876333	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 350 of SEQ ID NO:1432, b is an integer of 15 to 364, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1432, and where b is greater than or equal to a + 14.	", R23184, R68106, H04104, R7 D56912, D56797, F01477, R231 R68150, AI699279, R70543, R8 R48545, D56817, AI276541, AI R78505, R63400, AA459439, N2 M28696, M31933, X52473, M315 L08108 AA974370, W46279, AW196653 AA420451, AI29358, AA6690 AA521314, AA252310, AA2800 AI718165, AI129358, AA6690 AI718165, AI831132, AI027401 AI807828, Z40146, AA995204 AI807828, Z40146, AA995204 AI807829, AW369458, AA743770 D54675, AW149925, AW302960 AI340519, AI537677, AL1103 AI698391, AI929108, AW1292 AM059828, AW269098, AW2682 AM059828, AM349933, AL0364 AM300782, AI349933, AL0364 AIS81033, AI523989, AL1198
			s, Awsu0889, 5, ALO36396, 0, AW020397, 7, AW161579,
			AI813914, ALO80046, AW0895 AI753683, AW074993, AL0799

AL040169, AW268253, AI500659, AI950892, AI312152, AI815232, AI500523, AI468872, AW160916, AW162071, AI349937, AL036638,	7, AI345180, AW150578, AI62546 5, AL047042, AI252414, AW08040	AI633125, AW087445, AW071417, AI864836,	, AL037582, AL037602, , AI521012, AI312428,	, AW163554, AL135022, , AI610645, AI539771,	AL038605, AI343112, AW302992, Z99428, AI866770,	, AL121014, AI567582,	AI801325, AI815855,	AL119748, AL036980, AI889189, AL134830, AI890507, AW068845, AI612885, AA579618,	AI866820, AI564719,	AI497733, AL121365,	, AI091468, AI500662,	, AI472536, AW022808,	, AA493647, AI538850,	AI860783, AI624293,	AL036146, AL039/16, AW0/4869, AL30/543, AL047100, AI335426, AI348777, AW071362,	AL037030, AI569583, AI475371, AI635492,	6, AW075207, AI673363, AI34303	_	1, AI682841, AI859991, AL12069	, AA580663, AI568114,	7, AI683395, AL040456,	, AL036631,	AA641818, AC002350, AL096744, I48979, U35846,	L122050, I09499, I48978, Y166	AL110196, AL117457, U87620, AF090903, Y11587,

	AF090901, AL122093, AF100931, AL137550, M27260,
	AF090900, A08916, AL133606, AF078844, AL137538,
	A08910, AL049382, AF146568, AF090934, A65340,
	AL137271, AF183393, S78214, AL133565, A77033,
	Ħ
	AF113019, AL133557, I89931, AL050149, AF113013,
	X70685, AF
	0, AF079763, AL137533
	AL137557, AF031147
	AL117435,
	۲
	AF087943, AL049452, AL110221, X63574, AL050277,
	AL133080, I33392,
	AL137459, L31396, AL137527, AL050393, L31397,
	AF017437, AF118064,
	I49625, AF111849, S
	AR038854, AF118090
	AF057300, AF057299
	AL050172, Z82022,
-	, A03736, AF032666, A93016, AL
	AF111851,
	AF061943, AL133067,
	1, AL137479, U72620, AF113689
	AR059958, AF106697, U80742,
	×
	37294, S61953, AF118094, AL117583
	6558, AC
	D83032, AF100781, X80340, AF210052, A18777,

				AL117649, AL080158, 83508, AL137526, X875 U96683, AL137658, AL AL050155, A21103, AO L137292, AF081197, AL I17544, AF090886, Y1 L110218, AF119337, E0 67958, AF065135, AL13 AL050092, AJ012755, L049464, AL117585, AL L137463, AL137429, AR
1433	HOENU48	876334	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 2579 of SEQ ID NO:1433, b is an integer of 15 to 2593, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1433, and where b is greater than or equal to a + 14.	AA521311, AA521314, AW300598, AI051218, AIG31949, AA669095, AW298550, AA278335, AI694270, AW339489, AI797687, AA464762, AI948608, AI807828, AA810071, AA804200, AI718165, AA662808, AA504439, AI129358, AI718165, AA662808, AA504439, AI129358, AI718165, AA662808, AA50439, AI129358, AI718165, AA61280, AI299255, AA452985, AI718165, AA114888, AI129632, AA973497, N69756, T71487, AA252310, AA877638, AI027401, AA2555623, AA863081, AW196653, H47827, AA832206, AA995204, AA252340, Z28882, W46278, T48511, Z40146, AI831132, AA743770, D57019, AA344612, T84473, N87679, AI918466, Z19443, F00129, D56990, AI351209, AL047889, AW369458, AL047888, AC002350, D82786
1434	HOUDK26	876335	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 1038 of	H20994, H45211, H45368, H40040, H45293, H45192, AA205743, T24020, T90417, H20955, R70326, AF075043, AC004755, AC005516, AC005519, AL049836, AL080243, AC007358, AC005089, AC008394, AC00534, AC007546, AC005089, AL031597, AL031056, AC003690, AC005523,

			SEQ ID NO:1434, b is an integer of	ACOUZSIB, ACOU4881, ACOUZ472, H3U373
			correspond to the positions of	
		-	nucleotide residues shown in SEQ ID	
			than or equal to a + 14.	
1435	HODDG78	876340	Preferably excluded from the	AW247764, AA442668, AA491177, AW248120,
_			present invention are one or more	AL048314, AA479828, AA421873, AW248094, H75462,
			polynucleotides comprising a	Z42343, F06148, AA923747, F06007, AI445056,
			nucleotide sequence described by	R14715, F13060, AR025386, X86779
			the general formula of a-b, where a	
			is any integer between 1 to 651 of	
			SEQ ID NO:1435, b is an integer of	
			15 to 665, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEO ID	
			NO:1435, and where b is greater	
			than or equal to a + 14.	
1436	HAMFP80	876345	Ω	AI219740, AI478566, AI632246, AA279757,
			present invention are one or more	AA977612, AA716656, AA687260, AI801069,
			\vdash	AA071046, AI985849, AW370598, AA630617,
			nucleotide sequence described by	AW370599, AW370625, AA134295, AW390691,
			the general formula of a-b, where a	AI990289, AA134294, AA428452, AI143764, D30955,
			is any integer between 1 to 1090 of	AW370620, AA352142, AA074442, T83462, AW071043,
_			SEQ ID NO:1436, b is an integer of	T79236, AI744728
			15 to 1104, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1436, and where b is greater	
			than or equal to a + 14.	
1437	HWHQB10	876354	Preferably excluded from the	H40868
		•	present invention are one or more	
			polynucleotides comprising a	
			nucleotide sequence described by	
			the general formula of a-b, where a	
			is any integer between 1 to 345 of	

			SEQ ID NO:1437, b is an integer of				
			15 to 359, where both a and b				
			correspond to the positions of				
			de residues show				
			NO:1437, and where b is greater				
			than or equal to a + 14.				
1438	H2LAB47	876361	Preferably excluded from the	AA307985,	AL044985,	AA361756,	AA016093,
			present invention are one or more	AA133547,	AA046950,	AF126424,	AF106065,
			polynucleotides comprising a	AF076838,	AL122068,	AJ001642,	AJ131295,
			nucleotide sequence described by	AJ004977,	AF017748,	AF098534,	AF085736,
			the general formula of a-b, where a	AF106066,	AC004993,	AF098533	
			is any integer between 1 to 395 of				
			SEQ ID NO:1438, b is an integer of				
			15 to 409, where both a and b				
			correspond to the positions of				
	_		nucleotide residues shown in SEQ ID				
			NO:1438, and where b is greater				
			equal to a + 14.				
1439	HJBAR28	876364	Preferably excluded from the	AA355924,	N83684, A	AA214701, H	H94179, AW298728,
			present invention are one or more	AI056829,	AA278566,	AA093069,	AA278566, AA093069, T67190, AF092563
			polynucleotides comprising a				
			nucleotide sequence described by				
			the general formula of a-b, where a				
			is any integer between 1 to 390 of				
		-	SEQ ID NO:1439, b is an integer of				
			15 to 404, where both a and b				
			correspond to the positions of				
			nucleotide residues shown in SEQ ID				
			NO:1439, and where b is greater				
			than or equal to a + 14.				
1440	HCEFA76	876370	Preferably excluded from the	AL079827,	AA503895,	AB002353	
			present invention are one or more				
			polynucleotides comprising a				
			nucleotide sequence described by			,	
			the general formula of a-b, where a				
			is any integer between 1 to 338 of				

			SEQ ID NO:1440, b is an integer of 15 to 352, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1440, and where b is greater		
1441	нсовізі	876372	اعدا	7, AA446825, Z42384, W86347,	AC002064,
			present invention are one or more polynucleotides comprising a	T73581, T73682, T89320, T89957, R27248, R48643, H84547, H99963, N28347, N63131,	, R27450, , N64745,
			nucleotide sequence described by	A047464, AA047398, AA086034,	AA099567,
			the general formula of a-b, where a	AA099657, AA165569, AA169522, AA169441,	
			04.5 Qer	AA251391.	
				, AA258704, AA258149,	
			correspond to the positions of	0, AA622286,	
			nucleotide residues shown in SEQ ID	AA683138, AA713685, AA743062, AA807661,	
			NO:1441, and where b is greater	AA828448, D78955,	N87351,
	-		than or equal to a + 14.	AA165525, AA210972, AA211395, AA416558,	
				AA845854, AA971491, AA985073, AI023629,	
				6, AI092089,	
				, Z41403, Z45751, AI302012, AI3	57671,
				, AI367710, AI2017	
				AI445483, AI433348, AI478813, AI146981,	
				AI151439, AI184769, AI658554, AI521058,	
				AI537563, AI301471, AI634487	
1442	HTEGD78	876374	Ω		
	-		present invention are one or more	4I672898, AI874058, AI758608, AL079276	
					_
			nucleotide sequence described by		
			the general formula of a-b, where a		
	_		is any integer between 1 to 554 of		
			SEQ ID NO:1442, b is an integer of		
			15 to 568, where both a and b		
			correspond to the positions of		_
	•		nucleotide residues shown in SEQ ID		
			NO:1442, and where b is greater		

		qual to a + 14.	
1443 HCYBN59	876376		7, D80212, D80248, D80268, C14331,
		(1	D80227, D59927, D80269, D80133,
		polynucieotides comprising a	D80240, D803/8, D80166, D80219, D81
		nucleotide sequence described by	, C14389, D80157, D81030, C14429, D
		eral formula of a-b, wher	, AA305409, AW178983, D80195, D51060,
			2, D80366, D59859, D59502, D51423, D5179
		SEQ ID NO:1443, b is an integer of	D80045, D59467, C14014,
		15 to 654, where both a and b	D80391, D80164, D59787, D59275, D80043,
		correspond to the positions of	AA514186, D59889, D59610, D80193, D80196,
		nucleotide residues shown in SEQ ID	
_		NO:1443, and where b is greater	AW377671, AA305578, D59373, D80038, D80302,
		than or equal to a + 14.	AA514188, D80241, AW360811, D80247, AW177440,
····			
			C75259, D80258, AW178906, AW179328, AW366296,
			C05695, AW360844, AW360817, AW375406, D51103,
-			₹*
			AW178905, AW377676, AW378532, C06015, D80132,
			AW177501, DS9653, AW177511,
			W352171, AA
			D58253, D51250, AW17
			, AW178909, AW177456,
			3, AW178754, AW1790
			AW352117, AW178774
), D51213, AW179004, C03092,
			AW179012,
			D59695, AW378543, AI525923, AW352174, AW177728,
			N66429, AW
			AW367950, AW178911, AW177722, AI910186,
			AW378540, H67866, C14077, T11417, AW178781,
			AW360855, C14227, D58101, D51221, T03116,
			AW178986, AW177497, T02974, Z21582, AI535850,

			D59317, D59474, AW177723, D45273, C14973,
			AW3785
-			AI535686, C14298, AI55
_			551, AI525235,
	-		25215, AI525227, AI525912, AW37853
			42, AA285331, D50981, AW17901
			78542, AI525925, AI525222, C05763,
			955,
			-
			859
			2, A25909, A94995, Y12724, D89785, A7
			ω,
			3, D88547, AR066488, AR016514
-			X82626, A26615, AR052274, A43192, A43190,
			9, I82448, I14842,
	·		X68127, AR025207, AR054175
			, AR008281
			AR008408, AR062872, AR016691,
	-		A70867, A64136, A68321, D13509, AR060133,
			117, 179511
			5, T56234, T65208, R26874, R49147, R491
			R63286, R68208, R68209, R769
			N23372, N32910,
	_		W00634, W469
			5, AA046699,
			, AA131696, AA131540,
			ω,
			AA741529, AA767851, AA808213, AA812138,
•			2, AA938741,
			AA779560, AA868920,
			9, AI023812, AI093513,
			09719, AI274698, AI28
_			, AI478311, AI540692,
+			
1444 HCYBC31	876379	Preferably excluded from the	AA305023, AI352123, AI245481, AI909228, AI915162

	E C	т О	AC000402, AC002322
present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 885 of SEQ ID NO:1444, b is an integer of 15 to 899, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1444, and where b is greater than or equal to a + 14.	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 351 of SEQ ID NO:1445, b is an integer of 15 to 365, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1445, and where b is greater than or equal to a + 14.	preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 362 of SEQ ID NO:1446, b is an integer of 15 to 376, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1446, and where b is greater than or equal to a + 14.	Preferably excluded from the
	876380	876381	876382
	HCQBM44	HKCSP75	HKCSP84
	1445	1446	1447

			nresent invention are one or more	
			nucleotide seguence described by	
			the general formula of a-b, where a	
			is any integer between 1 to 289 of	
.			15 to 303, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1447, and where b is greater	
			than or equal to a + 14.	
1448 H	HPMFF45	876383	Preferably excluded from the	R52326, AL110125
			present invention are one or more	
-			polynucleotides comprising a	
	•		nucleotide sequence described by	
<u> </u>				
			is any integer between 1 to 511 of	
			SEQ ID NO:1448, b is an integer of	
			15 to 525, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
_			NO:1448, and where b is greater	
			than or equal to a + 14.	
1449 H	HE2CT52	876385	Preferably excluded from the	H74219, AA315682, AA904381
			present invention are one or more	
			polynucleotides comprising a	-
			nucleotide sequence described by	
			the general formula of a-b, where a	-
			is any integer between 1 to 605 of	
			SEQ ID NO:1449, b is an integer of	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1449, and where b is greater	
			than or equal to a + 14.	
1450 H	HTNBJ76	876386	Preferably excluded from the	AW083135, AA808057, AI745495, AA599616, T36219,

	present invention are one or more	AI918013, AA937922, A	_
-	polynucleotides comprising a	AI041990, AA342254, T	
	nucleotide sequence described by	AI360911, R11202, D25	5779, AI521589, AA076707,
	-	A1978792, AW068394, P	AA347093, AA323085,
	eger between 1 to 302		F17700, AL045709, AA077776,
	SEQ ID NO:1450, b is an integer of	AI633427, AA533408, P	AA558298, AA835710,
	15 to 316, where both a and b	AA330573, R87547, AI151261,	.51261, AI370475, AA297968,
	correspond to the positions of	AI699060, AI114477, 7	T92957, AI952780, AA972238,
	nucleotide residues shown in SEQ ID	AA857296, AA663306, V	W23546, AW268277, AA643261,
	드	l, AL042113,	F26719, AA825357, AI132963,
	equal to a + 14.	T47739, AI538812, AA548087,	48087, AA425924, AI890385,
		5, AI538540,	AA828762, H05073, AW419262,
		AW193493, AA527730, F	AI865988, T78484, AA468051,
		AW272763, AI049996, F	AI801141, AI913324, N84161,
		R82388, H82895, AW451	AW451360, AI053786, AI148927,
		2, AI0423	AA487219, AA384039,
		AA572960, AL046782, F	AA487079, AI754013,
		AA492313, AI923011, C	C13960, AW271904, AI753951,
		AA634209, AI755085, P	AA614010, AA235575,
	-	_	AI791150, AI623899,
		AA063139, AI114752, #	
		AA935377, AI859946, B	H73174, AA775049, AA581914,
		AI634323, AI470956, 7	_
			AA708678, AA311071,
		_	AI696901, AI754923,
		_	_
		AI755236, AI475332, 1	AL120976, AI915081,
		AA569182, AA664135, 1	AA831904, AA526656,
		AW189278, AA569743, 1	AA632845, AA714956,
		AA664789, AA525209, 1	AA507625, AI252506, Z36239,
		AI241705, AA776552, B	æ
		AI261913, AI275742, 1	AA82903
		, AC004253,	0
		AC007055, AL031055, 7	
		AL035587, AP000355, 1	AC005341, AL021391,
		AL049780, AC005209, 1	AL035455, AL034379,

	AL035450,	AL121655, U76377, AF029750, Z82172,
	AL109827,	AC005184, AC005778, AC006958,
	AC005071,	AL031257, AC009286, AC006132, Z82214,
	AL035687,	AC006146, AC004993, AL031295,
	AL049611,	AF001549, AC006115, AC005670, Z98257,
	AC004815,	AL121748, AL121603, Z85986, AL034421,
	S	AC007860,
	Z97200, A	AC002073, AL031767, AC004837, AC005666,
	AF196969,	AC005339, AC005011, AL035458,
	AF111169,	AC004797, AC005800, AL031846,
	AL121652,	AP000459, AL024498, AC006160,
	AC002045,	AC002472, AC002558, AC004485,
	AC005225,	AF190465, AP000112, AC006501,
	AC005624,	AC005081, AC005726, AC006026,
	AP000513,	
	AL031659,	AL050307, 297630, AL031054, AC004821,
	AC007406,	
	AC005088,	AL109967, AC007437, AP000036,
	AC007536,	AC007899, AC007114, AF042090,
	AC005480,	AC006547, AC004386, AC004876,
	AC005251,	1, AL022316
	AL080242,	AC006965, AC
	AF134726,	AC006013, AC006064, AL096774,
	AB020866,	, AP000211,
	AF064863,	, AL031311,
	AL035697,	AC005231
•	AL034547,	', L44140, AL021546, A
	AF146367,	Z98036, AP000144, AL031282, Z99128,
	AF053356,	AL133243, AL035451, AC007283,
	AC002996,	AC005082, AC010582, AL031589,
	AL034420,	AP001054, AL132985, AL034451,
	AC006116,	AF118808, AC006380, AC007298,
	AP000065,	AC002316, AP000088, AC005786,
	AC000003,	AC005598, AC005663, AC006978,
	73	C004050, AC002538, AC005284
	AD000216	793241 ACO07227 ALO49845 ACO04849

				AP000474	AC006344	275744 AC	AC007390. AL	AL049795.
				AI,022721	U91321. AC			AC010197.
				AP000517,		σ	9	
				AL031681,	AC003982,	AC005874,	AF134471,	
				AL132712,	AC004647,	AL078593,	AC007565,	
				AC005751,	AL031594,	Z82206, AI	Z82206, AL031286, AP00095	,6560000
				AC004000,	AC007510,	AC006530,	AC005280,	
		_		AC007649,	AP000230,	AC005971,	AC006480,	
				AL022165,	AC002364,	AL132992,	AC006323,	
		_		AC004020,	AC005821,	AF006501,		AC005799,
		_		AL050312,	AF038458,	AL021397,	U95742, AL	AL031121,
		_		AF124523,	AC004227,	AC003101,	AL022323,	_
		_		AF019413,	AJ229043,	AJ003147,	AP001037,	
_		_		AC006285,	AC009464,	AC006039,	AC005048,	-
		_		AC002377,	AP000692,	AC005245,	AC006597,	
		_		AC002365,	AL049643,	AL050318,	AC005057,	
		_		AC002115,	AC007221,	AC004814,	AC004111,	AL035462
1451	HE9ND38	876387	Preferably excluded from the	AA334551,	AA307537,	AF002996		
			present invention are one or more					
			polynucleotides comprising a					
			nucleotide sequence described by					
			the general formula of a-b, where a					
			is any integer between 1 to 351 of					
			SEQ ID NO:1451, b is an integer of					
			15 to 365, where both a and b					
			correspond to the positions of					
			nucleotide residues shown in SEQ ID					
			NO:1451, and where b is greater					
			than or equal to a + 14.					
1452	HPIAK40	876395	Preferably excluded from the	AI902815,		AI902293,	AR062079,	E05133,
			present invention are one or more				_	A27627,
			polynucleotides comprising a	E05329, E	E03742, E06(E06073, I19413,		E15669,
			nucleotide sequence described by	AR028747,	AR028747, A58083, E17345, I12374	17345, I12.	374, AR062080	080,
			the general formula of a-b, where a		E17344, E051			E05134,
			is any integer between 1 to 756 of	IS7961, E	E05162, E013	E01336, I12376	6, E17339,	E17340,
			SEQ ID NO:1452, b is an integer of	E17341, E	E17342, A37179	179, E05144	4, E05135,	121469,

			15 to 770, where both a and b	E05152, E05153, I21461, I90026, E05143, A14547,
			v	I21454, I31067
		·	nucleotide residues shown in SEQ ID	
			than or equal to a + 14.	
1453	HHPGD10	876397	Preferably excluded from the	AW361614, AB023235
			present invention are one or more	
			polynucleotides comprising a	
	•		nucleotide sequence described by	
			the general formula of a-b, where a	
	-	_	is any integer between 1 to 548 of	
			SEQ ID NO:1453, b is an integer of	
			15 to 562, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1453, and where b is greater	
			equal to a + 14.	
1454	HCQBI47	876398		AA527356, AI093930, AI635756, AW150892,
		_	present invention are one or more	AW340249, AI683004, AA574295, AA578334
			polynucleotides comprising a	
		·-	nucleotide sequence described by	
			the general formula of a-b, where a	
			is any integer between 1 to 1753 of	
			SEQ ID NO:1454, b is an integer of	
			15 to 1767, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1454, and where b is greater	
			than or equal to a + 14.	
1455	HE8DW67	876399	Preferably excluded from the	AA308646
			present invention are one or more	
			polynucleotides comprising a	
			nucleotide sequence described by	
			the general formula of a-b, where a	
			teger between 1 to 386	
			SEQ ID NO:1455, b is an integer of	

			15 to 400, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
	•		NO:1455, and where b is greater	
			than or equal to a + 14.	
1456	HONAH83	876400	Preferably excluded from the	AA398365,
			present invention are one or more	AA403200, N44265, AA362919, AI914181
			polynucleotides comprising a	
			nucleotide sequence described by	
			the general formula of a-b, where a	
			is any integer between 1 to 998 of	
			SEQ ID NO:1456, b is an integer of	
			15 to 1012, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1456, and where b is greater	
			than or equal to a + 14.	
1457	HHGCW95	876401	Preferably excluded from the	AA573757, AA161293, AA524449, AI742214,
			present invention are one or more	AA622626, W96506, AI476586, W96473, AA570007,
			polynucleotides comprising a	
			nucleotide sequence described by	AA502262, AI911816, AI796804, AA480659,
			the general formula of a-b, where a	AA552367, AI709265, AI809403, AI445236, AA552072
			is any integer between 1 to 623 of	
			SEQ ID NO:1457, b is an integer of	
			15 to 637, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1457, and where b is greater	
			than or equal to a + 14.	
1458	HCYB175	876402	பப	AA305438, AA056382, AW188096, AA308744,
			present invention are one or more	AI702438, C14389, D59927, C14331, D80022,
			polynucleotides comprising a	D50995, D80166, D80212, D80391, AW178983,
			nucleotide sequence described by	D59787, D59619, D80210, D80240, D80045, D80268,
			the general formula of a-b, where a	D58283, D81030, D80196, D59467, D51022, D59859,
			is any integer between 1 to 528 of	D51799, D80227, D80195, D51423, D80164, D59275,
			SEQ ID NO:1458, b is an integer of	D80253, D80043, D59502, AA305409, D80219,

	15 to 542, where both a and b	D80269, D80248, D81026, D80366, D80188, D50979,
	correspond to the positions of	2, C14429, C15076,
	ω.), D80193, D57483,
	_	0133, D80024, AA514188, AA514186,
	than or equal to a + 14.	AW360811, AW177440, D802
		178893, T03269, AW
		AW375405, D80157, AW178906, AW179328, AW366296,
		360844, AW360817, AW375406,
		, DS1759,
		AA056479, AW178905, AW378532, C0601
		AW352170, AW177501, AW177511, D51250, C05695,
		10132, AW
		AW178907, T48593, AW378528, AW178762, AW179019,
		AW179024, D80134, D59653, D58253, AW176467,
		1367967,
		AW179020, AW178775, AW178909, AW177456,
		AW360834, AW179329, AW178980, AW178914,
		AW178908, AW178754,
		, AW352117,
		3647, AW35
		AW378525, AW352163, T11417, D80949, H67854,
_		ည
		AW177728, AW367950, AA809122, AW179009,
		6, D80228, AI525923
		, AI905856, C14227, D45273,
		T03116, AI525917,
		, U88897,
		AC005145, AC004768, AL139054, AC005090,
		, AC006364, AC00720
		AF058696, A84916, A62300, A62298, AB028859,
		ò
		i, AB002449, X67155, A25909, AC00
		188, A94995, Y12724, A67220, D8
		D34614, AR008443, I50126, I50132, I50128,

				I50133, A43192, A4	A43190, AR060138, D88547,
				ω,	
				AR052274, I82448,	AR038669, X82626, Y09669
1459	HCRMK04	876404	Preferably excluded from the	AI057537, AI862687,	
			present invention are one or more	AA875951, AI783596	
			polynucleotides comprising a	AI374905, AI224513	
			nucleotide sequence described by	AI610450, AI829581	., AA775736, AI364904,
			the general formula of a-b, where a	AI698790, AA844090,), R71519, AI860091, AI523843,
			is any integer between 1 to 517 of	AI767012, AI473515,	3, AI350561, AW188551,
			SEQ ID NO:1459, b is an integer of	AL119399, Z99396,	Z99396, AL119324, AL119457, AL119443,
		_		AL042544, AL134524,	
			correspond to the positions of	AW392670, AL037051,	., AL036725, AA631969,
				AW372827, AL039074	i, AW384394, AL119497,
			NO:1459, and where b is greater	AL119418, AL036858	3, AL134920, AW363220,
			equal to a + 14.	AL036924, AL119483	3, U46341, AL119319, AL038509,
				AL039564, AL039085,	5, AL119396, AL039156,
				AL039108, AL039109,	
				AL119363, AL119341	., AL119391, AL119355,
				AL119335, U46350,	U46350, AL119522, U46349, U46351,
				AL119496, AL037094,	
				AL036196, AL036190,), AL037639, AL042965,
				AL038531, U46347,	U46347, AL042614, AL037085, AL119444,
				AL036767, U46346,	AL037082, AL042975, AL119464,
				AL037205, AL119488,	3, AL134533, AL119439,
				AL036268, AL039625,	
					3, AL03623
				AL042984, U46345,	AL038447, AL042909, AL039678,
				AL039629, AL134527,	7, AL042433, AL039386,
		·		AL042551, AL134531	1, AL039423, AL037077,
				AL042970, AL043029), AL042450, AL043011,
				AL043019, AL037615	5, AL038851, AL042542,
				AL036998, AL036733	_
				AL036765, AL036719	_
				AL036679, AL036774	, AL037021, AL036191
				4	
				AR069079, AB026436	5, AR054110

1460	H2CBF13	876405	Preferably excluded from the	AA307313, AA312913, AI203434
			present invention are one or more	
			polynucleotides comprising a	
			nucleotide sequence described by	
			the general formula of a-b, where a	
			is any integer between 1 to 593 of	
			SEQ ID NO:1460, b is an integer of	
			15 to 607, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1460, and where b is greater	
			than or equal to a + 14.	
1461	HKCSO44	876408	Preferably excluded from the	
			present invention are one or more	
			polynucleotides comprising a	
			nucleotide sequence described by	
			the general formula of a-b, where a	
			is any integer between 1 to 107 of	
			SEQ ID NO:1461, b is an integer of	
			15 to 121, where both a and b	
			correspond to the positions of	
			·O	-
			NO:1461, and where b is greater	
			than or equal to a + 14.	
1462	HWLKU83	876409	Preferably excluded from the	AW014464, AA693558, N74561, AI024015, AA332850
			present invention are one or more	
			polynucleotides comprising a	
			nucleotide sequence described by	
			the general formula of a-b, where a	
			is any integer between 1 to 692 of	
			SEQ ID NO:1462, b is an integer of	
			15 to 706, where both a and b	
-			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1462, and where b is greater	
			than or equal to a + 14.	

1463	HE9RM22	876418	Preferably excluded from the	AI492422, AI357898, AW296940, AA931635,
			ب ەب	
			nucleotide sequence described by	H81425,
			the general formula of a-b, where a	Z42746, Z42275, T89377
			teger between 1 to 175	
			SEQ ID NO:1463, b is an integer of	
			15 to 1765, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1463, and where b is greater	
			than or equal to a + 14.	
1464	HCRPQ93	876419	Preferably excluded from the	
			present invention are one or more	
			polynucleotides comprising a	
			nucleotide sequence described by	
			the general formula of a-b, where a	
			is any integer between 1 to 461 of	
			SEQ ID NO:1464, b is an integer of	
			15 to 475, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1464, and where b is greater	
			than or equal to a + 14.	
1465	HPDDL36	876420	Preferably excluded from the	AA366524
			present invention are one or more	
			polynucleotides comprising a	
			nucleotide sequence described by	
			the general formula of a-b, where a	
			is any integer between 1 to 184 of	
			SEQ ID NO:1465, b is an integer of	
			15 to 198, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1465, and where b is greater	
			than or equal to a + 14.	

1466	H2CBM09	876422	Preferably excluded from the	AA307727, AL121460, Z56847, Z57345
			present invention are one or more	
			polynucleotides comprising a	
			nucleotide sequence described by	
			whe	
			is any integer between 1 to 500 of	
			SEQ ID NO:1466, b is an integer of	
			15 to 514, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1466, and where b is greater	
			than or equal to a + 14.	
1467	HKCAA10	876425	Preferably excluded from the	AA192455, AW294111, AA707196, AI924499
			present invention are one or more	
			polynucleotides comprising a	
			nucleotide sequence described by	
			the general formula of a-b, where a	
	-		is any integer between 1 to 635 of	
			SEQ ID NO:1467, b is an integer of	
			15 to 649, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1467, and where b is greater	
			equal to a + 14.	
1468	H2CB125	876426	Preferably excluded from the	AA307505, AA360083
	-		present invention are one or more	
			polynucleotides comprising a	
			nucleotide sequence described by	
			the general formula of a-b, where a	
_			is any integer between 1 to 465 of	
		-	SEQ ID NO:1468, b is an integer of	
			15 to 479, where both a and b	
			correspond to the positions of	
_				
			NO:1468, and where b is greater	
			than or equal to a + 14.	

1469	HKISB80	876427	d by	AA718982
			is any integer between 1 to 385 of SEQ ID NO:1469, b is an integer of 15 to 399, where both a and b correspond to the positions of	
			residues shown ir d where b is greadle to a + 14.	
1470	H2CBE84	876428	oly excluded from the	AW009512, AI609285,
•			present invention are one or more polynucleotides comprising a	AA301898, AI671626, AI818892, AW025713, AA490857, R40307, AA700491, AI273067, AA834371,
			nucleotide sequence described by	AI368173, AW316631, C05075, AA480122, AA348046,
			mula of a-b, wher	, AA089704, D80241, D59467, Z21582
			IS any inceger becween I to 44% of SEO ID NO:1470, b is an integer of	DS8283, D81039, D59839, D51423, D80188, D80188, D80188, D58283, D58283, D81030, D59619, D80210, D51799, D80240,
			15 to 460, where both a and b	, D59889, D80195, D80038, D80022,
			correspond to the positions of	D80043, D80391, D59275, D57483, D59787, D80227,
			nucleotide residues shown in SEQ ID	2, D80366, D80196, D50995, C14331,
			NO:1470, and where b is greater	9, D50979,
			than or equal to a + 14.	, C14014, C15076, AA305405
				, D58253, C04935, AW178893
				D51022, D80949.
		,		32, AW177440, AA30
		•		111, D80251
				AW178762, AA514188, C14298, D80133, AA514186,
•				C14407, AW360811, AI557751, AW378540, D51097,
				C05695, AW375405, AW360834, AA285331, AW377671,
				«366296, АW360817, AW
				AW179024, D80302, D59373, AW179020, AW177456,
				AW352171, AW377676, AW178906, AW352170,

			AW177731, AW178907, AW178754, AW179019, D80247,
			9004, AW179012, D517
-			AW179018, T11417, H67866, T03116, D80157,
			AW178781, AW378525, D51103, I
			557774, AW352120, AW17728, AW178
			AW178911, AW378543, AW352163, D80258, D59653,
			D45260, T02974, D59503, D51213, T48593, H67854,
	-		
			AW367950, D80064, AW178986, AI525923, D58246,
	. .		05
			C14973, AJ525920, C14046, D60010, AI535686,
			AI525912, AI525227, AI525215, AC002036, A62298,
			AJ132110, A84916, A62300, AR018138, D88547,
			D34614, X67155, Y17188, D89785, D26022, A25909,
-			A67220, A78862, AR008278, A45456, X82626,
			AF058696, AB028859, AR025207, Y12724, AB012117,
			X68127, AR066482, A85396, A82595, A44171,
	-		A85477, A94995, I19525, A86792, U87250,
			X93549,
			', I50126,
			R066488,
			4, Y09669,
			A63261, D88507, AR
-			U46128, AR008408
1471 HSEBD08	3D08 876431	Preferably excluded from t	, AW242810, AI888669,
		present invention are one or more	, AA773636,
		ides comp	AA699864, AA112388,
			AI524767, AW377081, AW016549, D62897, AA954644,
		the general formula of a-b, where a	AA169505, AW377047, AA092662, AW362046,
		teger between 1 to 1993	AA629163, S72869
		SEQ ID NO:1471, b is an integer of	
		15 to 2007, where both a and b	

	R42236, AI268027	AI913961, AA621915, AI768685, AW009951	AI744435, AA725348, AI910436, AA771917, AW275132, AI915670, AI217575, AA772389
correspond to the positions of nucleotide residues shown in SEQ ID NO:1471, and where b is greater than or equal to a + 14.	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 386 of SEQ ID NO:1472, b is an integer of 15 to 400, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1472, and where b is greater than or equal to a + 14.	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 1264 of SEQ ID NO:1473, b is an integer of 15 to 1278, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1473, and where b is greater than or equal to a + 14.	preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 461 of SEQ ID NO:1474, b is an integer of 15 to 475, where both a and b
	876432	876435	876436
	HPMFM22	нонев 14	HAIDH43
	1472	1473	1474

			correspond to the positions of nucleotide residues shown in SEQ ID NO:1477, and where b is greater than or equal to a + 14.		
1478	HE8UJ03	876447	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 2757 of SEQ ID NO:1478, b is an integer of 15 to 2771, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1478, and where b is greater than or equal to a + 14.	AW340972, AI763378, AI745530, AI400359, AA634799, AW373755, AA406542, AW008882, AI379597, AW373615, AI858439, AI380423, AI628029, AW074041, AI538874, AW189012, AA857364, D82303, AA224830, AA132792, AA224831, AA524982, AW364047, AI678604, AI142902, AA133068, D82445, H39906, AA593133, AA644624, AA133068, D82445, H39906, AA593133, AA644624, AA888921, AA411736, AI992380, AI679729, AA904079, AA494400, AA577041, AI282492, AI640743, AW074288, AI535647, AA551421, AA336073, AA505483, AI469669, AI284099, AI284098, AI201463, AI903549, AI903561,	59, 32, 12, 12, 74644624, 29, 29, 29, 72,
1479	HDTLK03	876448	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 2051 of SEQ ID NO:1479, b is an integer of 15 to 2065, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1479, and where b is greater than or equal to a + 14.	340 526, AW391549, 160, AA085664, 338, AA984772, 348, W26560, AI 348, AA581646, 643, AI335437, 074, AW089030, 002, AA648105, 680, AA334191, 719, N71529, AA 820, D20893, AI	AW304931, AA659697, N22162, AA085613, 311237, AI336661, AI344929, AA847210, AI382955, AI933533, AW370221, 186588, AW363311, 186588, AW363311, 3116, AI251367,
1480	HMTBC69	876451	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by	D50810, U62768, U62769, U32990, U76997 AJ131025, AJ131026, AJ131027, AJ131028	97, 28

			the general formula of a-b, where a is any integer between 1 to 706 of SEQ ID NO:1480, b is an integer of 15 to 720, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1480, and where b is greater than or equal to a + 14.	
	HMUBP81	876452	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 1153 of SEQ ID NO:1481, b is an integer of 15 to 1167, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1481, and where b is greater than or equal to a + 14.	H 4 4 4 1
1482	HAPOT58	876458	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 2115 of SEQ ID NO:1482, b is an integer of 15 to 2129, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1482, and where b is greater than or equal to a + 14.	AL037788, AI686047, AI753484, AI636777, AI861877, AI935355, AI144560, AI192999, AI806026, AA081086, AI140416, N52261, AI984946, AI126835, AI375382, N31999, AI431922, AI000687, AA281546, AI354844, AW368199, AI806020, AI192995, AA432212, AI796776, AI765555, AI436119, N62465, AA416953, AI392798, AA504837, AA993835, AI342228, N74643, AA962052, N31979, H80204, AI340563, AW025654, W95677, AI373352, AA958965, AA505730, AA598619, AA281547, AA455805, AI350119, AI143974, AA283875, AI810436, AI761126, AA456624, AA931610, AI634994, AI149059, H58033, AA282093, AI762032, AI867892, W39405, W15216, AA456424, AI493979, W26521, AI418808, W95891, AA470851, N92893

				H81006. AA136357, AA359333, N50738, AI309586,
				3, AW293385, AA373138
	_			
				AI589997, AA605260, AA370986, AI690377,
				AA359446, W73659, H78829, AA113788, AI761221,
				AI469943, AA609846, AI864350, W25612, R24652,
				AA360514, AI907228, AA831054, AA355628, H78428,
				AI473940, AA291183, AA745877, AA136269, T24969,
				AI693730, AA706077, N83393, AA070852, AI905829,
				AI587625, N88059, AW363223, AI559993, AA526788,
				AI216608, AW371352, AI634388, N79184, AW363222,
				AA594328, AA400847, AI209205, AA393670, H83189,
				AF161432
1483	HCFLR18	876459	Preferably excluded from the	AA807288, AL036653, AL036654, AI289925, AI291875
			present invention are one or more	
			polynucleotides comprising a	
			nucleotide sequence described by	
			the general formula of a-b, where a	
			is any integer between 1 to 519 of	
			SEQ ID NO:1483, b is an integer of	
			15 to 533, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1483, and where b is greater	
			than or equal to a + 14.	
1484	HDPAA38	876464	Preferably excluded from the	AA873176, AA931378, AI218111, AI014843,
			present invention are one or more	AC005379
			9	AL096702, AF187320, AL117258, U95740, AC004797,
			nucleotide sequence described by	295704, AC004636, AC005071, AP000952
			the general formula of a-b, where a	
			is any integer between 1 to 887 of	
			SEQ ID NO:1484, b is an integer of	
			15 to 901, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
	:		NO:1484, and where b is greater	

			than or equal to a + 14.	
1485	HCYBM66	876465	erably excluded fraction are nucleotides comprised by incleotides comprised in the position of the position of the position are nucleotide residues shall be sequence despend to the position are nucleotides comprised in the position of the position and where position and and and and and and and and and an	AA116082, AA305687, C14014, D80269, D80227, AA809122, AA305409, C14389, D80391, D59787, D80196, D58283, D59859, D80022, C14331, D80166, D80195, D59467, D51423, D59619, D80210, D51799, D80164, D59275, D80240, D81030, D80253, D80043, D59502, D80212, D80188, C15076, D80219, D59927, D59502, D80024, D59610, D80378, H67854, T03269, C14429, AW178893, D80241, D80045, AW178532, AW36951, AA305578, AW178775, AW178762, D51250, AW360511, AA305578, AW178775, AW178762, D51250, AW360811, F13647, D80522, C14227, D58253, AW360811, F33647, D80522, C14227, D58253, AW360811, AA31618, AB377671, AW377676, AW360834, AW366296, C05695, AW352171, AW360844, D81111, AW360817, AW377672, AW179023, AW178905, D80064, D80268, C14298, AW378534, AW178905, D80064, D80268, C14298, AW17731, AW178907, AW178906, AW352170, AW177731, AW178907, AW178906, AW352170, AW177731, AW178907, AW178908, AC07649
1487	HLTAH77	876470	equal to a + 14. ly excluded from	AI359524, AW003850, AI089719, AI359474,

			nresent invention are one or more	AI652055, AI948841, AI824819, R87348, F13369,
			Ä	ın
			nucleotide sequence described by	F08357, AF035282
			the general formula of a-b, where a	
			, -	
			SEQ ID NO:1487, b is an integer of	
			15 to 1181, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1487, and where b is greater	
			than or equal to a + 14.	
1488	HWLXX39	876471	Preferably excluded from the	AI879483, AA553761, AW363300, AW162358
			present invention are one or more	
			polynucleotides comprising a	
			nucleotide sequence described by	•
			the general formula of a-b, where a	
			is any integer between 1 to 491 of	
		_	SEQ ID NO:1488, b is an integer of	
			15 to 505, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1488, and where b is greater	
			than or equal to a + 14.	
1489	HPTWG85	876472	Preferably excluded from the	AI652564, Y17108, Z92544, Y17258
			present invention are one or more	
			polynucleotides comprising a	
			nucleotide sequence described by	
			the general formula of a-b, where a	
			is any integer between 1 to 637 of	
			SEQ ID NO:1489, b is an integer of	
			15 to 651, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1489, and where b is greater	
			than or equal to a + 14.	
1490	HE6BS09	876473	Preferably excluded from the	AL120741, AA573741, AW409804, AA191552, W93042,

present invention are one or more	AW402618, AW409704, AA496304, AW073345,
 polynucleotides comprising a	AW300845, AA744892, N39760, AW176264, AI498051,
nucleotide sequence described by	, AA5048
 the general formula of a-b, where a	AI564499, AI128977, AA737814, AA419313,
is any integer between 1 to 2954 of	, N26317, AW
 SEQ ID NO:1490, b is an integer of	
15 to 2968, where both a and b	AA307525, AI272853,
 correspond to the positions of	N35109, AA191421, AI091816, W24942, N62754,
nucleotide residues shown in SEQ ID	1, R35445,
NO:1490, and where b is greater	4, AA8642
 than or equal to a + 14.	AI219732, R75982, AA506884, AA868134, N95815,
	7, AI2771
	AA55250
	72,
	T78250, AL07
	R27494, AA348004,
	AA337541, AA356674, T48679, AA73837
	\sim
	5658, AI20
	`
	1364021,
	AL079579, AA665375, R79989, AA355436, R34256,
	AA368982, AA348005, AA327401, N43853, AA937676,
	ò
	AA337180, AI520916, AI684053, AA054425,
	AI866770, AA878790, AI890907, AI348854,
	AI608932, AW001426, AI358701, AI680498,
	AIS54343, AI620639, AL038445, AI961589,
 	AI758437, AA911767, AI611348, AW022682,
	AW131288, AA603709, AI288285, AI344935,
	AI310575, AL037582, AL037602, AI340533,
	AL042191, AI349645, AW268253, AI702301,
	AI345253, AW083175, AI349937, AI621209,
 	AI345026, AI559531, AI554485, AW150804,
	40627, AI9638
	AI335235, AA908294, AW105601, AI497733,

AI340511,	AI263331,	AL036980, AT.036904.
45739, AI340659,	AI343091,	AI251221,
9, AI932638,	719	81
AW268072,	AI335208,	AW089275,
4, AW079336,	588	3 6
, AW074993,	AW129106,	AI445131,
AI349957,	AI348777,	AW301300,
۷,	AI312152,	AI343037,
AI470293,	AI889148,	AW152469,
, AI349226,	AI345005,	AI348879,
, AI886206,	AW058233,	AI349614,
2, AW193134,	AI307543,	AI307210,
	AI349598,	AW302988,
3, AI349256,	AW167222,	AI313320,
_	AI620284,	AW023338,
_	AI307520,	AW088805,
_	AI334884,	AW071412,
, AI312325,	AI343140,	AI349971,
, AW081	AI783504,	U49908, M30514,
E02349, I48978, AL117	435, X849	
9, AL0494		X83508, AL049314,
A08916,	1, A08913	I03321, A08913, AJ238278,
A08909,	943, AR02	X6357
0	AL137521, AL133568,	33568, I49625,
AL050393, AJ012755, AR038854, AF028823	R038854,	AF028823,
	11880, AF	R038969, AL133016,
4	0, AF1136	594, AF113690,
	F158248,	AL023657, AF158248, A18777, AF079763,
AL117457, E02221, X53587, AF090896,	587, AF09	90896, AF118094,
	AF113677,	
	AL137292,	E04233, U58996,
, AF017437,		0, I4240
, AL050116,	92,	, AF00843
AL050172, Y10080, AL1	AL110197, AE	AF111849, AL117649,

			AF090900, AF125949, S78214, AF061943, I26207,
			18631, U7
			2, S75997, AF091084, AL049452, X7
-			6, AF113019,
			I96214, AF215669, AL137478, AL110196, AL110280,
			A07647, AL137558, AL050138, AL133072, AL137480,
			L049300, AI
			AL133081, Z37987, AF162270, U00763, AL137429,
			I09360, AL050024, AL080124, AL133098, AL117460,
			AL096744, AF026816
			on
			2889, AL13
			774
			526, AL137523, AF097996
			836
			Y09972, AL050108, AL137488, AF106657, AL133113,
			AL080234
			ς.
			77033, A77035, AF087943,
			AF111851, AL110221,
			676, I66342, AL137533,
			AL122121, AF032666
			AL122049,
			. 7
), Z82022
			L31396, U68387, AL133077, AF177401, S68736,
			5, AF0909
			E08264, E07361, A93016, S61953, A21103,
			37459, X00861, AF1262
			7582, AL122050, AL137529,
			12297, AF057300, AF057299, AL110171, AL08006
\dashv			A08907, AF113689, AF017152, AL133075, AR068751
1491 HERAM35	876474	Preferably excluded from the	

			present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 515 of SEQ ID NO:1491, b is an integer of 15 to 529, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1491, and where b is greater than or equal to a + 14.	
1492	HFIUG54	876475	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 1211 of SEQ ID NO:1492, b is an integer of 15 to 1225, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1492, and where b is greater than or equal to a + 14.	AIO96476, AI627324, AW176260, AA420479, AI888162, AW001768, AW166776, AIO17162, AIO86151, N20484, AA AA126992, AW370989, N34406, AW391594, AA N24599, CO2570, AW38 W105105, AA570014, AW AW020880, Z41211, AI H24299, AA678544, AW AA613111, AI925770, D51223, D62210, AA84 N75648, AI436629, N5
1493	HE8CX56	876476	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 2284 of SEQ ID NO:1493, b is an integer of 15 to 2298, where both a and b correspond to the positions of	AI693062, AI936680, AI638780, AW130947, AI203659, AA969048, AA730307, D61225, AL041011, R49279, H64578, AA249856, AA120957, H64682, D81623, AL040722, N56191, AW265781, AA082593, AF029343

SEQ ID	AW068683, AA314376, D80193, D80210, D80240, D59467, D8010 D80210, D80240, D59467, D8010 D80164, D59275, D80038, D8026 D51423, C14331, D59859, D8026 D50391, D80253, D81030, D50505 D505051, D80378, D80241, D80378, D80378, D80241, D803740, D51022, AW352158, D80134, D51250, D52291, AA51AW178762, AW1777501, F13647, D80134, D51250, D52291, AA51AW366296, D80132, AW360817, AW366296, D80132, AW360817, AW366296, D80132, AW360817, AW366296, D80132, AW3777731 AW179023, AW377676, AW179018 D80247, AW378528, AW178908, T11417, A1557751, AW178914, AW378525, D51103, AW178914, D80157, AW378533, D59627, C060 AW378533, A1557774, D4525920, D52417, AM179013, D80122, C03092, H67854, D8AM179013, D59217, AN525920, D5AM179013, D59217, AM525920, D5AM179013, D59217, AM525920, D5AM179013, D59217, AM525920, D5AM179013, D59217, AM525920, D5AM179013, D592020, D5AM179013, D59200, D5200, D5200, D520020044, D59200, D50200, D50200, D50200, D5020044, D50200, D502	
nucleotide residues shown in SEQ NO:1493, and where b is greater than or equal to a + 14.	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where is any integer between 1 to 375 of SEQ ID NO:1494, b is an integer of 15 to 389, where both a and b correspond to the positions of nucleotide residues shown in SEQ IN NO:1494, and where b is greater than or equal to a + 14.	
	876480	
	H2LAQ54	
	1494	

				J132110, A62300, A622 67155, D26022, A25909 34614, D88547, AF0586 B028859, AR025207, Y1 R066482, A94995, X681 AR008443, A85396, A4 19525, A86792, I50126 93549, AR066488, AR01 26615, AR052274, Y096 43190, AR038669, I183 188507, I14842, AR0541 63261, AF135125, AR00 A70867, AR008277, AR U46128, D13509, A641 AB033111, AR064240
1495	HWABG32	876481	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 1386 of SEQ ID NO:1495, b is an integer of 15 to 1400, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1495, and where b is greater than or equal to a + 14.	AA873178, AW340076, AA453258, AA453359, AI200335, AI189856, AI127354, T57079, AA031327, AI096450, AA948375, AA031328, AA977624, AA994405, AI148795, AI340956, AW014990, AI652909, AI160243, AW026239, AI093526, AA923811, AI091630, AI365268, AW380222, AA579988, AI367151, N32402, AA583097, N56822, AA579988, AI343747, H12681, AI825678, AW197534, T29148, F08275, AI468467, T95661, T82166, T57151, AI880292, T81821, F04505, AA481266, R41605, AW372903, AA662708, AW130992, AI818777, AA764938, X14356, L03418, X14355, L03419, M91645, M91646, M91647, M82819, L03420, M63835, M91555, M91552, AI81820, AI81830, S45707, AA109725
1496	HMTBE05	876483	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by	AI026945, AI808573, AI620239, AA948677, N53940, AW249558, AI096948, AA159915, AI095014, AI871045, AI950931, AA455901, AW009419, AI149374, AA024477, AI433743, AA428948,

			the general formula of a-b, where a	AA039950,	AA165025,	AI884373,	AI149074,	
			teger between	AI184801,	AI188603,	AI937231,		
			SEQ ID NO:1496, b is an integer of	AI469664,	W26293, A	W26293, AA831823, AI766893,		AA830218,
			15 to 1484, where both a and b	AA476574,		AW404545, AA45590	AA455902,	-
			correspond to the positions of	AA027936,	AI566799	AA582203, R15907,	R15907, AA	AA422121,
			nucleotide residues shown in SEQ ID	AI879131,	T34650,	43817, AA7	Z43817, AA738453, AI220916,	20916,
			NO:1496, and where b is greater			AI300117, AA738075,	A738075, AI	AI967928,
			than or equal to a + 14.	Z39886, A	AW071642, A	AA863299, A		AI382238,
				AI149361,	AW169605,	AA483840,	AI436690,	•
				AA448896,	AI800263,	AI831898,	AI262999,	
		_		AI984945,	AI915652,	AI701265,	AI344209,	M79093,
				AI829004,	AA028041,	AW408623,	AI982982,	
				AI202924,	AW246104,	T66533		
1497	HKABL05	876484	Preferably excluded from the	AI740522,	AI309318,	AI376662,	AI741390,	
			present invention are one or more	AI742840,	AA679083,	AI765150,	AW002945,	
			polynucleotides comprising a	AW192895,	AA001262,	AI052703,	AA648295,	
			nucleotide sequence described by	AI929375,	AW157334,	AI799150,	AA577690,	
			the general formula of a-b, where a	AA909347,	AA608744,	AI879998,	AI421323,	WS5919,
				AW373539,	W84527, AA947742,		AA861283, AA	AA065133,
			SEQ ID NO:1497, b is an integer of	AW168112,	AA460061,	AI300565,	AW204198,	
			15 to 2192, where both a and b	AA155821,	AW104051,	AI800773,	AI193965,	
				AA101195,	AI582368,	AW057835,	AI348116,	
			nucleotide residues shown in SEQ ID	AA527861,	AW009823,	AW029295,	AW022530,	
	,		NO:1497, and where b is greater	AA708118,	AW238854,	AI452699,	AI016610,	
		_	than or equal to a + 14.	AA669337,	AA480279,	AA278360,	AI749692,	
				AI160871,	AW130090,	AA744919,	AA760760,	
		_		AW007135,	AI275625,	AI057288,	AI494111,	
				AA831711,	AA687284,	AI815697,	AI374689,	
				AA155925,	AI862854,	W55920, AI367891,		W04222,
				AW272692,	AA628638,	AA707011,	AI800064,	
				AA043251,	AA160009,	N62094, A	AI671739, AA	AA292750,
				AI052618,	AW166814,	AA152365,	AI475145,	N78325,
				AA001852,	AI952464,	AI953334,	AI346774,	
				AI243902,	AI271553,	AI637742,	AA514862,	
				AA025382,	AA484277,	AI288842,	AI311020,	NS0975,
				AW027908,	AA132226,	AI436690,	AI130684,	N74257,

A	2, AI3	54226, AI96	9402,	AI026752,
A)	AA453035, AA6	AA668696, AI	AI090673,	AA971631,
N N N N N N N N N N N N N N N N N N N	AA984913, AW26		AI798057, I	N93127, AL120009,
A PA			ò	AW163390,
A			AA764824,	AI521457,
AI	AW439109, AA08	AA088421, AA	AA722831, 1	N23855, AA807549,
- F	AA043590, W678	307, AA02	6016, AA	×
 - Pi	38,	AA065202, AA	AA928577,	AA633795, W15314,
A.	AI886794, W84!	W84515, AI79	7422, AA	AI797422, AA120907, AA046354,
A)	, 76	AA083453, AA		AA009957,
A.	AI190992, AA20	AA284411, AA	AA857371,	AA459969,
 A	AA741542, AA00	_	AI206746,	AA160010,
A	AA586336, AW235	920,	AA010759,	AW075660,
- A	AA131616, AA0	AA046070, AA	AA247207,	AA002267,
 AI	AW020230, AII:	AI123351, AA	AA281235,	AA426610,
 - Ai	AA780786, AI82	5394,	AA083357,	W73815, AI439077,
A	AI434359, AI695	507,	AI344209,	W69764, W60465,
A:	AI281441, AA5	AA568376, T6	3795, W3	T63795, W38654, AA028052,
A.	AI826611, AI80	AI800263, AW	AW270667, AI370333	AI370333,
Al	AW117628, W52	113, AA12	7865, AW	W52413, AA127865, AW439098, T64108,
P	AA164988, AA2	11263, AA	278324,	AA211263, AA278324, AA327661, C15972,
\$	W78007, AA011120,	120, T470	65, T348	A20
A		N66464, AA49	AA491375, AA	ο,
P			D54180, AA	, AI984
A.	, 57	AW341620, AW	AW438482, N99121	, AA054
 <u>A</u>	AA226936, T943	Ø	6323, AA	227046, AI559910,
- A	AA574112, AI2	AI290025, AA		AA460014,
A PI	AW050391, AA9		AA373413,	AA356295,
 <u>~</u>	AA621388, AWD	AW009092, AA	301008,	AA301008, AA482700, T64028,
- F	AA332547, T35		5052, T6	AA205052, T63820, AA738461,
A	AI000546, N33	N33952, T570	17, AI88	T57017, AI887555, AA365643,
 - A	_	28948, AA	448896,	AA428948, AA448896, AA211143, T51962,
<u>ex</u>	R15907, AA131382,		AA142894, T30133,	0133, AB030905,
 À	841, Z8			AF
A .	AB005618, X56	X56683, A75245,		75, D28
<u>5</u>	U09120, AF086	16270, T47064	64, T52042	42 R16239

				N38911, N	N46485, N58965, W39742, AA028051,
					30, AA147058.
				AA186506,	AA278348,
				AA525773,	AA525871, AA661828, N56031, C00146,
				AA091857,	
				AA665715,	, AA732979, Z18797, AA99182
				AI001836,	Z39146, AI341188, AI566368, AI652212
1498	HOCTA74	876487	Preferably excluded from the	AI302800,	AW118693, AI808667, AI065036,
			present invention are one or more	AW080952,	AA862461, AI201847, AI138543,
			polynucleotides comprising a	AI015998,	AA865819, AA470462, AA454546,
			nucleotide sequence described by	AI221895,	AA481881, AI039771, AA535254,
			the general formula of a-b, where a	AA482063,	AI301489, AA551867, AI018725,
				AL121442,	AI244932, T88913, AI914566, AI017732,
	**************************************		SEQ ID NO:1498, b is an integer of	AI016693,	
			15 to 685, where both a and b	AA120922,	N57711, AW151576, AI572464, AW303732,
			correspond to the positions of	AI471156,	R85699, H60433, AA890675, AI262997,
			nucleotide residues shown in SEQ ID	AA620388,	T47276, AA534566, AI625454, AA852619,
			NO:1498, and where b is greater	AA889211,	AA707578, AI718799, T47275, AI124998,
			than or equal to a + 14.	AA477467,	H88225, AA680222, H66348, N63309,
				AA131070,	AA131015, AI474581, AI561334,
				AW392670,	AW372827, AW
				AL119497,	AL134528, AL119443, U46341, AL119457,
				AL119319,	
				AL119324,	AL119355, AL119483, AL119484,
				AL119391,	AL042965, AL119335, U46350, AL134920,
					AL119396, U46351, U46349, AL119418,
					AL119444, U46346, AL037205, AL134902,
				AL042614,	
				AL042551,	AL119401, AL134518, AL134524,
				AL043029,	AI142132, U46345, AL042984, AL134531,
				AL134538,	AL134525, AL042450, AL043019,
				AL134536,	AL037051, AL036725, AL042970,
				AL119488,	AL042544, AL042542, AL043003,
				AL119464,	
				AL122056,	A81671, AR066494, AR060234, AR054110,
				AB026436,	AR069079

HWLUU48	876490	preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 1035 of SEC ID NO:1499, h is an integer of	AA099027, AI566740, AW085264, AI383310, AA453266, AL020989,	AI887335, AW086500, AI590636, AA436251, AC004190, AC007100	AI887905, AI222690, AA411391, AI913708, AP000516,	AI694672, AI686357, AI431702, AI015064, AB014087,	·
		•					
HULA115	876491	Preferably excluded from the	AI991884,	AI872008,	AI660228,	AW167205,	
		invention are one c	AW084525,	AA601542,	AI859727,	AI818462,	
		polynucleorides comprising a	AWUSUSS,	A168/516,	AA35611,	AMODO876,	
		the general formula of a-b, where a	AI569542,	AI860861,	AI887280,	AI653757,	
		is any integer between 1 to 1004 of	AA461121,	AI554798,	AI016349,	AA622753,	
		SEQ ID NO:1500, b is an integer of	AI332503,	AI246460,	AI332793,	AI144192,	
		15 to 1018, where both a and b	AA460819,		AA455216,	AA621675,	
		correspond to the positions of	AA862530,	AA858222,	AA581826,	AI806046,	N35715,
		nucleotide residues shown in SEQ ID	AW328329,	AI262551,	AI204029,	AI149450,	
		NO:1500, and where b is greater	AW071084,	AI289219,	AA609900,	AA927266,	
		than or equal to a + 14.	AI707484,	AI095745,	AA618130,	AI721109,	
			AA931503,	AI440027,	AI275080,	AI299248,	
			AI276688,	AI750085,	AA088417,	AA304654,	
	*****		AI262552,	AI688181,	AI282807,	AW294666,	
			AI335810,	898	AI335786,	AA088540,	
			AA420995,	AI355863,	AA102237,	AA070673,	
			AA595597,	AI750051,	AI749025,	AI811127,	
			AI086655,	AI278320,	AA443973,	AI080248,	
			AI367574,	AA421075,	AA052939,	AI418137,	
			AA902863,	AI265947,	AA931116,	AA430411,	
			AA251968,	AI355088,	AI290353,	AW305028,	
			AI005354,	AI367787,	AA913300,	AA053492,	
			AW008828,	AI355089,	AI890124,	AA564009,	

				AI359453, AI282383, W45582, W52209, AA102236,
				9, AI0951
				S
				AI70806
				AI720469, AA879062, AA186928, AA494466,
				AI832504, H79930, AA417983, W45545, AA469124,
				AA526593, AI719480, AI832612, AA420865,
				AI264706, AA242885, N35628, AA858264, H62987,
	·			, AA865264, AA418153
				AA353482, AA740793, AI310701, AI143647,
				AA320588, AI541426, AI581554, AA420466,
				AI472533, AA188357, AI888688, AA373467,
				8, T61575,
	-			
				AI582088, W79666, AA377021, W74128, AA370626,
				AA126713, R23407, U46351, AA193598, AI581181,
				9, T61023, H96296,
				, AI123178,
	-			, AA576977
	_			HS
	-			, M57710,
				X78879, U06470, X16834, J02962, J03723, X16074,
	-			
	_			24, AF0314
┥	_			A59344, M27260, AL122093, AL117599, AL133015
1501	HSYAJ64 8	876494	Preferably excluded from the	
			present invention are one or more	_
	-		polynucleotides comprising a	AW377280, AA581816, AI435156, AA599212,
			mula of a-b,	AI828823, AA160573, AA894927, AA446427,
-			between 1 to 2017	, AA314621, AA812415,
			SEQ ID NO:1501, b is an integer of	AA307680, AW377313, AA315193, AA514946,

	, wh	AA948141,	AA652118,	AI090292,	AA43551,	
	correspond to the positions of	AI342258,	AI240388,	AA205318,	AA243054,	
	res	AA768432,	AI082283,	AA024693,	AA456625,	
	NO:1501, and where b is greater	AI911813,	AI363735,	AA446119,	AA652124,	
	than or equal to a + 14.	AA424926,	AI263712,	AA024647,	AA205575,	
	•	AI004571,	AA630601,	AA307175,	AA164747,	
		AI042562,	AI934643,	AI341665,	AA313490,	N75485,
		AA207213,	W91894, A	AA426166, AA307366,		AI433060,
		AA307046,	AA195483,	AA252561,	AA527990,	*
		AA989506,	AA223574,	AI270387,	AA243053,	
		AA455806,	AA307677,	AW403863,	AA315014,	
		AA159366,	AA157555,	AA158206,	AI568188,	
		AI028221,	AI445024,	AA927196,	AA307925,	
		AA649534,	T28878, A	I085919, A	T28878, AI085919, AW392054, AA776680	776680,
		AI672839,	AA312108,	AA376260,	AW392206,	
		AA654257,	AI865398,	AA347324,	AA626750,	
		AA219493,	AI630717,	AA307419,	AA662020,	
-		AI510831,	AA442877,	AA350306,	AA362375,	
		AI935046,	AA152328,	AI305172,	W05296, AI	AI278536,
		AI308922,	AA053461,	AA053213,	AA135056,	
		AA186979,	AW173202,	AW377352,	AA206750,	-
		AA608732,	AI025236,	AI719108,	AA325720,	
		AI922470,	AA223615,	AA152329,	AA626448,	
		AA649822,	AA300684,	AA362586,	AA626522,	
		AW377293,	AA315660,		AA333552, R3	R37150,
		R15974, A	AIS69355, A	AA190772, A	AA362376, AA	AA593069,
		AA921347,	AA316929,	AA180011,	AA134971,	W95113,
		AA978212,	AI932667,	AA040890,	AA830424,	
-		AW383641,	AI632334,	AA947203,	AA326527,	
		AA629781,	AW383640,	AA954366,	R05778, C2	C21408,
		R05864, A	W392327, A	AW392327, AA191382, AA32273	ις ,	H55311,
		AW383658,	R15975,	AW410508, A	AA995270, AA	AA160528,
		AA219455,		AI703040, AW104153,	M27396,	M15798,
		M27838, X	52130, U07	M27838, X52130, U07201, U07202, U38940	2, U38940,	
		AC005326,	L35946, M	AC005326, L35946, M27054, L35936	936, L35937	
		L35938, L	35945, L35	L35945, L35940, L35941,	L35942,	L35939,

				7.35943 1,35944 1,35935	TERRON TERRO1
COST	TETIELO	0.104.00	December 2001 1. 200 1. 100	OCCUR ALONIN AGAACOK	100001 (200001
7001	HELLFLY	α/σ υν υν	റ്	AA326636, HI68/4, AW3/6009, AA313468, K23401.	009, AA313468, K23401.,
			present invention are one or more	N35321, R13283, AW152493,	3, AI027550, T11328,
			polynucleotides comprising a	AR036119, X92689, U70538	8
			nucleotide sequence described by		
			the general formula of a-b, where a		
			is any integer between 1 to 1449 of		
			SEQ ID NO:1502, b is an integer of		
			15 to 1463, where both a and b		
			correspond to the positions of		
			nucleotide residues shown in SEQ ID		
			NO:1502, and where b is greater		
			equal to a + 14.		
1503	HLYEA23	876496	Preferably excluded from the	AW161801, NS6973, N73756,	6, AA479038, D44982,
			present invention are one or more	N81193, W65438, H25021,	W65438, H25021, N22293, N47355,
				AA973373, AA477521, AA5	AA595499, AA838190,
	_		nucleotide sequence described by	AW172858, AI887235, AL1	AL134275, T59612, AW169038,
	_		the general formula of a-b, where a	AA847980, AI002744, H02	H02058, AI590442, AB014528,
			is any integer between 1 to 556 of	AC005062, AL135783, AL1	AL117258, AL133163,
	_	-	SEQ ID NO:1503, b is an integer of	AL137100, AC004859, AL0	AL035410, AC004067,
			15 to 570, where both a and b	AC002349, AC005725, AF2	AF205588, AC008033,
			correspond to the positions of	AC004887, AL049589, AC0	AC002412, AF130249,
			nucleotide residues shown in SEQ ID	AC005261, AC007488, AL0	AL033533, AC005722,
			NO:1503, and where b is greater	AC007011, AC006547, AC0	AC006080, Z98304, Z84469,
			than or equal to a + 14.	AC005664, AF031078, AF0	AF030876, AF031076, 295152,
	_			AC004019, AC005280, Z69	Z69907, AC006213, AC007238,
	_			AL049569, Z93016, AP000344,	344, AL031597, AC004605,
				282203	
1504	HAPQU61	876498	Preferably excluded from the	AI949815, AI813450, AI8	AI819294, AI269353,
			present invention are one or more	AA421819, AI089074, AA8	AA834705, AA847960,
			polynucleotides comprising a	AI559836, D31784	
	_		nucleotide sequence described by		
	_		the general formula of a-b, where a		
			is any integer between 1 to 484 of		
	_		SEQ ID NO:1504, b is an integer of		
			15 to 498, where both a and b		

			correspond to the positions of nucleotide residues shown in SEQ ID NO:1504, and where b is greater	
1505	НЕ8ОТ93	876499		AA486504, AA133234, AI339710, AA743093, AI688621, AI096844, AA129712, AI860744, AI420708, AI278953, AI278568, AW006666, AI571986, N68247, AI358873, AA314945, AA341071, AI346152, AI219397, AA488692, AA148150, AI362046, AW050985, AI090396, R60368, AA626449, AW272569, AA308535, AI471517, AW135592, AW205875, R60312, AI590397, AI078709, N39886, AA557504, AA970783, AI419556, AA338145, AA557504, AA970783, AI419556, AA338145, AA530976, NS7132, AW051845, AW394065, H95626, AA309736, AW204673, AI457186, AA376417, AA570135, AI805191, AA376416, AA310109, N68052, H95981, AI049818, Z21567, AA079141, AW389275,
1506	H2LAB08	876503	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 2382 of SEQ ID NO:1506, b is an integer of 15 to 2396, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1506, and where b is greater than or equal to a + 14.	AI911983, AI927427, AI889004, AI693602, AL045565, AI767631, AI150323, AA576743, AI201732, AA811424, AA436321, AI890062, AA812674, AI348111, AA776471, AA904047, AI909133, AA262396, AI909125, AA827237, AW084600, AI890814, AA778086, AI708713, AA436197, AI580236, AA688044, AA252436, AA307642, AI569986, AI174417, AA251902, D19596, AA307642, AI569986, AI174417, AA251902, D19596, AA307642, AI569986, AI174417, AA251902, D19596, AA307642, AI66986, AI774417, AA251902, AI000199, AA307642, AI66986, AI174417, AA251902, AI6789252, AI67893, AI67832, AA651878, AA307939, AA378903, AI934157, AA243609, AI267661, AA525290, AI824311, R37260, R59445, AA378902, D61809, AA361618, R12332, AA831575, R75944,

), AA354320, AA602417, AI56
			N87729, H03382, H01205, R31246, H00817, R39541,
			R92975, AL045564, D58065, AA730991, C16596,
			v
			3
			AI289791, AI539800, AI500714, AI355008,
	_	-	AI866469, AI434242, AI539771, AI889189,
			_
-			AI582912, AI927233, AI433157, AI612913,
			AI491710, AI366900, AI804505, AI610362,
			AI434223, AL039390, AI440239, AI863197,
			AI924051, AI366910, AI539847, AI521596,
			AW074057, AI932620, AL040207, AI590043,
			AL042944, AI567935, AI539260, AI866465,
			AI801325, AI500523, AI538850, AI887775,
			AIS37187, AI923989, AI284517, AI872423,
			AI500706, AI445237, AI491776, AWI51138,
			AI521560, AI500662, AI284509, AW172723,
			AI440263, AI538885, AI889168, AI866573,
			,
			AI284513, AI888118, AI285439, AI859991,
			9, AI889147,
			_
			1, AI866786,
			6, AI242736, AI828574,
			73, AI539781, AI539707,
			AI885949, AI285419, AW089557, AI559957,
			AI521571, AI469775, AI866581, AI567953,
-			AI815150, AI446495, AI867068, AI225248,
			AI610426, AI567940, AI282264, AI926593,
			AF035293, AF081281, AF052112, AF077198,
			AF077199, D63885, AC004062, U97146, AR028701,
			U97147, U97148, U89352, AC004548, AL133074,
-			Y17793, AL133076
1507 HISBB72	876504	Preferably excluded from the	AI589824, AW149545, AA826266, AI285235,
		present invention are one or more	AA548396, AI580850, AI934791, AI262821,

				C03639, AA361522, AA370109, AW131681, T48460.
				AA350472,
1510	HWLQP42	876513	Preferably excluded from the	AA196276, AA524473, AL040260, AA533568,
			present invention are one or more	AA600703, AA773551, AA292150, AA004500,
			polynucleotides comprising a	AI928071, AI612760, AA411191, AW264086,
			nucleotide sequence described by	AW206769, AA496356, AA434061, W42808, AA232555,
			the general formula of a-b, where a	AI085934, AA182481,
			teger between 1 to 999	AA004501,
	-			AW317087, AI752948, AA443125, AA456190,
		-	15 to 1013, where both a and b	AA400594, AA292028, AI682335, R73572, AA766115,
	-		correspond to the positions of	AA292042, H61296, H61291, AL043495, AA044201,
			nucleotide residues shown in SEQ ID	R11520, AA705241, AA652065, AA043939, AI536587,
			NO:1510, and where b is greater	R97731, AI352191, AI630315, AA350112, D31167,
			than or equal to a + 14.	AA031359, T85323, AA429498, H15771, R44134,
-		_		AI351143, AW138388, AA661960, AI215409,
				AA411071, AW243696, R72952, AW068860, AI567210,
				AI393957, AI970891, AI273925, AA321611,
				AA401967, AI224608, AI084609, AI279699,
				AA031603, AI915877, AA400679, AI092030,
				AA031637, AI630462, AA429499, AA031476,
				AA301177, H15770, AW381505, AA182758, AW381475,
				R10445, AW381498, AI992085, AA312507
1511	HDPAG07	816918	Preferably excluded from the	AA305114, AL022398
	-		present invention are one or more	
			polynucleotides comprising a	
			nucleotide sequence described by	
			the general formula of a-b, where a	
			is any integer between 1 to 442 of	
			SEQ ID NO:1511, b is an integer of	
			15 to 456, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
		_	NO:1511, and where b is greater	
			than or equal to a + 14.	
1512	HLTAR39	876524	Preferably excluded from the	AI133655, T96748, AW369762, AA350015, AA360756,

			nresent invention are one or more	AW386072, AI625829, AA534216, AW243183,
			. a	AI697340, AI754731,
			nucleotide sequence described by	AC004707, AC004675, AF088219, AC006026
			the general formula of a-b, where a	
			.,	
			SEQ ID NO:1512, b is an integer of	
			15 to 2167, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1512, and where b is greater	
			than or equal to a + 14.	
1513	HWLRF38	876526	Preferably excluded from the	AW183028, N28485, AI306451, AI536589, AW072566,
			present invention are one or more	N24976, H82376, AI814709, AI376566, AI352453,
			polynucleotides comprising a	AI590303, AI280262, AI761747, AA554283,
			nucleotide sequence described by	AI222990, AA644328, AA661978, AA587549,
			the general formula of a-b, where a	AA045302, AW274520, AW043629, AA630727,
			is any integer between 1 to 818 of	AW273650, AI368900, AI381943, AI290422,
			SEQ ID NO:1513, b is an integer of	AI167243, AA993296, AA977315, AW337456,
			15 to 832, where both a and b	AA029935, AA779545, Z17865, AI493253, AI624318,
			correspond to the positions of	AA908755, AI168437, AA757538, AA977243,
		-	nucleotide residues shown in SEQ ID	AI740891, AA524068, AA628420, AI123070,
			NO:1513, and where b is greater	AI692442, AI868044, AA687907, AI370323, T31450,
			equal to a + 14.	AI867272, N46853, N67292, AW276010, N69329,
				AI768256, AI022628, R83171, AW073539, AA180796,
				AI761569, AA045408, AW134931, AW085513,
				AW059629, D11973, AL133563, AJ006412, AB018284,
				AJ006776
1514	HCRNM09	876530	Preferably excluded from the	AW362945, AI916280, AA632418, AW451840,
_			present invention are one or more	AA579245, R85405, AW366782
			polynucleotides comprising a	
			nucleotide sequence described by	
			the general formula of a-b, where a	
			is any integer between 1 to 1350 of	
			SEQ ID NO:1514, b is an integer of	
			15 to 1364, where both a and b	
			correspond to the positions of	

AW129659, AI250293, AI432969, AI445025,	AI868831, AW166645, AI500077, AI349645,	5628, AL049085, AI538829, AW15148	AW303152, AL045266, AL043326, AI567351,	AI345111, AW238730, AI619502, AW268253,	AL044207, AW026882,	, AI343112,	, AI783504, AI690426, AI6311	AL079963, AI866608, AL121014, AI857296,	AI499512, AI349004, AI281762, AI590227,	AI873731, AL038778, AW088793, AI440426,	AI580984, AI269862, AL119828, AI439762,	AI570909, AI673710, AI815232, AI340582,	AI682841, AW103371, AI635461, AW149869,	AI625079, AI274541, AI610756, AW008048,	AL043975, AW089572, AI348897, AI922901,	AI636456, AI538259, AI312152, AI282903,	AI349937, AW162071, AW129202, AW169132,	_	AI634737, AW090013, AI249257, AI636445,	4, AW196141,	AIS54484, AI811344, AI912866, AI570384,	AW002342, AI569616, AI475451, AI702433,	AI224992, AI799199, AI271786, AI273142,	AI432656, AA508692, AW068845, AI269696,	AI800433, AI560099, AW132121, AI284517,	AA613907, AI498579, AI445165, AL117613,	AF147302, AF090900, I48979, AF113694, AL080124,	AL133640, I89947, Y11587, AF090934, S78214,	AF113691, AL133606, AF090903, AL117460,	AL049938, L31396, AL122093, L31397, AF104032,	AF078844, AL050146	, AL117457, S68736, AF090901, AF11301	50393, AL122050, I89931, AL	AL050149, AF113690, AF113677, AF118070,

				AF090943, AL137459, AF113013, AL110196,
				AF113676, AJ242859,
<u> </u>				AF090896,
				38913, AF01
				AF113689, Y16645, AL049452, AR059958, AL096744,
				AL137527, AL133565, AL049314, AL122123,
				AL049466,
				AL133093, AF111851,
				AF091512, Y11254, AL117435,
-				AC002464,
				AC004686, U91329,
				AL133560, AL110280, AL117394, AL049430,
			•	AL110225, AJ012755, AC004383, A65341, AL078602,
				14
_				I66342, AF042090, AC006501, U95739, AL049382,
				E07108, AC007458,
		-		E02349, AL117585,
				U00763,
				C002467, AF
				AL031732, AC010077, AL122110
1516 HATCV09	.V09 876534		Preferably excluded from the	AI949332,
		pr	present invention are one or more	AW169558, AA857218, AI433853, AW204540, R68303,
-		<u>o</u>	polynucleotides comprising a	AA994295, AI
		nu	nucleotide sequence described by	R44174, Z40075, AI015727, N34408, R74002,
		th	the general formula of a-b, where a	R68268, R53421, R54010, Z38312, R44219, R49558,
		is	is any integer between 1 to 2095 of	AA090402, F01959, AA090979, U72788, AI304833
		SE	SEQ ID NO:1516, b is an integer of	
		15	15 to 2109, where both a and b	
		8	correspond to the positions of	
		nu	ide residues show	
		<u>8</u>	NO:1516, and where b is greater	

			than or equal to a + 14.	
1517	HCRNE16	876535	y excluded from nvention are one otides comprisir e sequence descral formula of a-teger between 1:1517, b is an i, where both a ed to the positic e residues showr and where b is gqual to a + 14.	, C06072, AI589250, AI470584, AA , AA747122, T27280, AC007501, U8
88.	HCRPV63	876536	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 411 of SEQ ID NO:1518, b is an integer of 15 to 425, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1518, and where b is greater than or equal to a + 14.	AI143683, AI924826, AA086365, AI792153, Z79581, Z79582, S81107
1519	HSKKP02	876538	preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 1172 of SEQ ID NO:1519, b is an integer of 15 to 1186, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1519, and where b is greater	AA916748, R83779, AA331626, AA400220

			than or emial to a ± 14	
1520	HOVANI3	876540	blv excluded f	
í)))	present invention are one or more	
			nucleotide sequence described by	
			=	
			integer between 1 to 446	
			SEQ ID NO:1520, b is an integer of	
			15 to 460, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
		,	NO:1520, and where b is greater	
			than or equal to a + 14.	
1521	HWBEX78	876543	Preferably excluded from the	W20138, AA229752, AI380196, N44538, AA026809,
			present invention are one or more	R41836, N71112, N33777, W05473, AA026870,
			polynucleotides comprising a	W15415, AA888089, W39614, R68936, AI143439,
			nucleotide sequence described by	H00351, R63287, T54159,
			the general formula of a-b, where a	W91983,
_			is any integer between 1 to 1658 of	
			SEQ ID NO:1521, b is an integer of	
			15 to 1672, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			than or equal to a + 14.	
1522	HRODG74	876544	Preferably excluded from the	AI797095, AA902901, N47240, AI252632, AI718169,
			present invention are one or more	AW079806, H09548, AI203811, AA459245, D25745,
			polynucleotides comprising a	C21350, R63205, AC006065, AC002368, AF025422
	_		nucleotide sequence described by	
			the general formula of a-b, where a	
_	_		is any integer between 1 to 574 of	
			SEQ ID NO:1522, b is an integer of	
		•	15 to 588, where both a and b	
			correspond to the positions of	
_			de residues s	
			NO:1522, and where b is greater	

			than or equal to a + 14.	
1523	HCROK30	876545	preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 506 of SEQ ID NO:1523, b is an integer of 15 to 520, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1523, and where b is greater than or equal to a + 14.	, AA682308, AI540716,
1524	HDABK73	8 7 8 7 9 9 9 9 9 9 9 9 9 9 9 9 9 9 9 9	preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 2777 of SEQ ID NO:1524, b is an integer of 15 to 2791, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1524, and where b is greater than or equal to a + 14.	A1744148, A1744113, A1860811, A1889014, A1765413, AW237314, A1765401, AL042645, A1867571, AW293518, AA534578, A1432178, AN867571, AW293518, AA534578, A143237, AA406169, AW188054, A1147954, AA430324, AL040186, A1197943, A1589634, AA569041, A1015938, AA4313904, AA070872, A1188829, A1124780; AA421239, A1149224, AA420647, A1916160, W73655, A1076564, A1768356, R51293, A158215, A125307, W51790, AA172002, AA425349, AA565222, AA313542, AA825728, R35270, AW204507, AA100809, W28763, A1222042, A1479185, W26572, W45413, W73608, R52192, A1160529, AW440819, A1422286, A1298011, AA171761, AA421279, R51403, H62930, R52097, R59309, AA581790, W81419, A1768849, W40121, A1708313, AA373236, AW368276, AA434583, Z42217, W81420, A1962360, AA325784, R59310, A1271621, T25845, T06069, F05246, AA806028, Z38264, AA071023, AA815452, N54389, AA810542, AA383377, A1370602, R50941, T87272, T87186, F01748, AA947741, AA773493, AA890049, A1985779, AA984284, AW272799, AL043147, AB007891
1525	HOGCO78	876548	Preferably excluded from the	AI471995, AW393929, AA044743, AI741975,

	present invention are one or more	AA044797,	AI720824,	AI992258,	AI480029,
	polynucleotides comprising a	AI803250,	AI095557,	AI245572,	AA662934,
	sednence des	AA876346,	AW327457,	AW393932,	AW157188,
	the general formula of a-b, where a	AI669783,	AI286104,	AA025525,	AI090194,
	eger between 1 to 673	AI128230,	AI095934,	AI189306,	AI950299,
	NO:1525, b is an	AI467898,	AA028934,	AI742307,	AA194396,
	15 to 687, where both a and b	AI809949,	AI160162,	AI122798,	AI034059,
	correspond to the positions of	AI244940,			AI431317, AA746600,
	le residue	AI150927,	R19215, A		R96173, AW043889,
	NO:1525, and where b is greater	AA876265,	AA844331,		4, AA860575,
	than or equal to a + 14.	AA487470,	AI432084,	U56654, A	AW157607, AA669015,
		AI825990,	AA335548,	4.	AA932576,
		2.	AI270663,	AI497894,	AI221399, R13183,
			AA564849, A	AI866853, A	AW272239, AW150208,
		AI572774,	AA668506,	r	AI866127,
		AI568138,	AA641818,	AI923370,	AW118518,
		AL038665,	AW264727,	AI582932,	AW078818,
		AI866469,	AI687168,	AL037582,	AL037602,
		AI241923,	AI613038,	AI473536,	AI866465,
		AI559872,	AI955117,	AW020095,	AW078606,
_		AI288285,	AW090451,	AL046942,	AW079409,
		AI635016,	AL079963,	AI827058,	AIS90043,
		vo	AI687166,	AI620302,	AI611738,
		AI446721,	AI961589,	AL041772,	AI500061,
		AI457589,	AISS9752,	AW166870,	AI125884,
		AI687127,	AI802542,	AI452707,	AI932503,
		AL039132,	AI581362,	AI624293,	AI434656,
-		AIS87279,	AI561228,	AW051226,	AI348870,
		AA983883,	AL135024,	AI289542,	AI554821,
		AI453339,	AL138420,	AW149925,	AW150557,
		AI915291,	AL039086,	AW163834,	AI654276,
		AW026882,	AI433157,	AW083572,	AI702073,
		AA225339,	AI860897,	AI418681,	AL036638,
		AI923989,	AI800341,	AW131294,	AI539800,
		AI621341,	12	AI698391,	AI538564,
		AL040827,	AL046466,	AW152182,	AI270429,

	AI355779, AI695726, AI638644, AI628325,	
	AI819014, AI818980, AW079075, AI357644,	
	AW262552, AI927256, AW128834, AL046595,	
	AI636588, AI651840, AW054964, AL119399,	
	AW264895, AI884318, AI889189, AL120995,	
	_	
	AI612913, AI469270, AW024793, AI818353,	
	AW105459, AI866770, AI445303, AI309306,	
	AI475806, AI267185, AI583558, AI932794,	
	AW410259, AI686576, AI335214, AW148294,	
-	AW198090, AI270706, AA502794, AL039716,	
	AI891084, AI520702, AI691088, AI569975,	
-	AI434731, AI538817, AI571439, AI279925,	
	_	
	AL036673, AI670002, AI335426, AI348777,	
	AW051088, AI819976, AI927233, AI912438,	
	AI491842, T69241, AI963846, AI873638, AI565172	65172,
	AW148544, AI270183, AI699823, AW263355,	
	AI612750, AI540674, AI817523, AW087915,	
	AL043152,	
	٥,	
	AI538885, W74529, AW	46618,
	AI929108, AI446373	
	, AW196078, AI67	37480,
	A77033, A77035, I89947, AL1220	
•	L133640, AF008439, AF111849,	47716,
	X63162, AF106657,	02578,
-	AL096744, A08910, A08909, AL	751,
	, A08908, AF090903, AR038854,	434,
	L137557,	80154,
	AL117457, A08913, Z97214, A65340, AF107847,	7,
	A08912, E06743, AF111112, I4	
	L117460, A0	, 8,
	AF026816, AF215669, AL133075,	U78525,
	13, ALO50092, AR034	
	AF061573, U58996, A58524, A58523, AF090934,	4,

AF113677, Y14314, AL050155, AL117435, S78214, A86558, AL049938, AL049466, AL137550, AL133014, AF090896, D83032, E05822, X84990, AF017437	L050172,	AFILSOLS, ACCES, ACCOSTOR), U68233, I92592, AJ003118	AL110280, U77594, I49625, A65341, AL050024,	AL137558, I32738, AF030513	A03736, L04504, U889	U42766, AF028823, AF113699, Y09972, AF124728,	AL023657, AL133665, AL080148, AL137521,	, AL122110, I89934,	AL133645, AF115392, Z13966,	A93350, S69510, AL137533, AF177401, AL117440,	AL050138, AL133010, AF182215, X83508, AF100931,	E02349, AF061981, AL137479, U72620, A15345,	AL137539, AF097996, AF067728, AL137478,	9, AR029490, AL133557, E01314,	AF118094, AF090943,	AL133016, A23630, AF081197, AF081195, AL117648,	56039, X6	L31396, S77771, AL137537, L1	AL049314, A49139, AF061795, AF151685, S83440,		U67958, Y10936, AL049430, X80340, AF118092,	7, AF017152, I09	AL133619,	5, Y10080,	AL133637,	, AF139986,	, AF091084, AF113690	0, E04233, AL080110, U49434, AF026	U96683, AL110221, AL117578, U87620, A58545,
5														-						-									

				D16301, AL137658, U72621, AL080126, AF104032,
	,			3. I68732, E12747, AL133560,
				AF078844, AL080060
1526	HCRNG10	876549	Preferably excluded from the	AA737831, AA651628, AI239587, AA912347
			present invention are one or more	
			polynucleotides comprising a	
			nucleotide sequence described by	
			the general formula of a-b, where a	
-			SEQ ID NO:1526, b is an integer of	
			15 to 708, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1526, and where b is greater	
			v	
1527	HWLRR08	876551	Preferably excluded from the	AI040700
			present invention are one or more	
	-		polynucleotides comprising a	
			nucleotide sequence described by	
	_		the general formula of a-b, where a	
			is any integer between 1 to 604 of	
			SEQ ID NO:1527, b is an integer of	
			15 to 618, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1527, and where b is greater	
			than or equal to a + 14.	
1528	HTEFPSS	876553	Preferably excluded from the	AA454500, AW301277,
			present invention are one or more	AW388282, AA129369
			polynucleotides comprising a	AW450017, AW418819, H56484, AA437031, AW082355,
			nucleotide sequence described by	AW204742, U28413
			the general formula of a-b, where a	
			is any integer between 1 to 1089 of	
-			SEQ ID NO:1528, b is an integer of	
			both a and	
			correspond to the positions of	

		AA934986, AA782950	AA490688, AW264544, AW378307, AA961504, AA613715, AW021810, AA554460, AW368463,
		AM315265,	AW271245, AI018136, AW384563, AW086214, AW020066, AW192488, AI783695, AW384566,
	·	AI797860, AA569967,	AW117930, AA315280, AW384544, AW384497, AW192483, AI917637, AA677120, AW378298, AI816732,
	AL110374	AI207993, AA621885,	AW276060, AI598114, AW378323, AW383155, AA257102, AA461400, AA315269, AI589498, AA461087,
nucleotide residues shown in SEQ ID NO:1528, and where b is greater than or equal to a + 14.	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 206 of SEQ ID NO:1529, b is an integer of 15 to 220, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1529, and where b is greater than or equal to a + 14.	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 424 of SEQ ID NO:1530, b is an integer of 15 to 438, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1530, and where b is greater than or equal to a + 14.	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 2048 of SEQ ID NO:1531, b is an integer of 15 to 2062, where both a and b correspond to the positions of
	876557	876558	8 7 6 5 5 9 5 5 5 9
	HDLAR46	H2CBW66	HOGDS65
	1529	1530	1531

			nucleotide residues shown in SEQ ID	AW368530, AI341438, AW378317, AI290266,
			NO:1531, and where b is greater than or equal to a + 14.	AW368521, AI280695, AW384490, AI418400, AI970613, AI160977, AW023591, AA947181,
				,
				N71882, AI
				AI952506, AAZ5/UI/, AA490466, H88912, N69323, AI912481, AA055599, N67469, M86849, I74304,
1532	H2CBX36	876560	Preferably excluded from the	
			present invention are one or more	M85168,
				AB035424, AB035422, AB035425, AB035423, AB035421
			nucleotide sequence described by	
			the general formula of a-b, where a	
			is any integer between 1 to 1144 of	
			SEQ ID NO:1532, b is an integer of	
			15 to 1158, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1532, and where b is greater	
			than or equal to a + 14.	
1533	HSHAX43	876572	Preferably excluded from the	H66220, AA809449
			present invention are one or more	
			polynucleotides comprising a	
			nucleotide sequence described by	
			when	
			is any integer between 1 to 562 of	
			:1533, b is an	
			15 to 576, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1533, and where b is greater	
			than or equal to a + 14.	
1534	HCRQI57	876575	Preferably excluded from the	AI361150, AI939490, AW089648, AF002993
			present invention are one or more	
			polynucleotides comprising a	
			nucleotide sequence described by	

				AW352158, AW178914, AW17733, AW178908,
				AW178754, AW179018, AW369651, AW352117, T48593,
				4, D45260, AW179012, AW178774, AW
				AW352163, AW352120, H67866, F13647, AI525923,
				80
		-		H67854, AW179011, AW179009
	-			, AI905856, D59503
				D58246,
•		_		AW178781, C13958, AI535686,
				1, C14407, AI525917, D45273, C1
				AA514184, AI525227, AW378533
				V178986, AI525920
				5, C14957, C14046,
	•			C14298, T03048,
				_
				D51097, C16955, T02868, U13896, U13897, U93309,
				U51639, A84916,
				A62300, A62298, AR018138, AR008278, AF058696,
				Y17188, D26022,
				IS0128, IS0133, D88547, I14842, AR066488,
			-	AR016514, X82626, AR060138, A45456, A26615,
_				
				\vdash
				AR008281, AR066490, AR062872, A70867, AR016691,
				٠.
	- , -			A64136, A68321, X68127, AB012117, D13509,
				AR060133, X72378, A85396, D88507, AR066482,
		_		AF123263, A44171, AR032065, A85477, I19525,
				A86792, X93549
1536	HHEGC16	876579	Preferably excluded from the	, AA573289,
			present invention are one or more	, AW372737, AW38
		-1	polynucleotides comprising a	AI951269, AI560208, AW372734, AI309528,

			nucleotide sequence described by	AW372745. AA121349. AI097133. AI310351	51,
			the general formula of a-b, where a	AW073286, AI160271,	01,
			ω ω	AW170797, AW388634, H69344, AA278853, AW372735	, AW372735,
			SEQ ID NO:1536, b is an integer of	H47623, AA742972, AA864447, N31288, AW372730,	AW372730,
			15 to 1532, where both a and b	AI572193, AA173309, AW188877, H69345,	, AW363751,
			correspond to the positions of	AW372731, AW372736, H47925, AI476011,	
			nucleotide residues shown in SEQ ID	AA278420, AW372739, AW372744, H38254,	, N22901,
			NO:1536, and where b is greater	AA278794, AA769896, AW372740, AW37278	86,
			than or equal to a + 14.	AW372738, AL040673, AF132937	
1537	H2CBG53	876580	Preferably excluded from the	AA307226, AB020236, AF045449	
			present invention are one or more		
			polynucleotides comprising a		
•			nucleotide sequence described by		
			the general formula of a-b, where a		
			is any integer between 1 to 468 of		
			SEQ ID NO:1537, b is an integer of		
			15 to 482, where both a and b		_
•			correspond to the positions of		
			nucleotide residues shown in SEQ ID		
			NO:1537, and where b is greater		
			than or equal to a + 14.		
1538	HCYBF23	876581	Preferably excluded from the	AI949966, AA687405,	50,
			present invention are one or more	AA721257, AW028336, AA305220, AI522235	35,
			polynucleotides comprising a	AA827201, AW298461, AI220695, AI984660	60,
			nucleotide sequence described by	AI219204, AI026116, M84722, M84721, D12775	D12775,
-			the general formula of a-b, where a	U90888,	10, D88988,
			is any integer between 1 to 709 of	D31634, U29907, D31637, U29911, D88989	68
			SEQ ID NO:1538, b is an integer of		
			15 to 723, where both a and b		
_			correspond to the positions of		
			nucleotide residues shown in SEQ ID		
			NO:1538, and where b is greater		
			than or equal to a + 14.		
1539	НОБСО80	876583	Preferably excluded from the	AW076027, R24903, R32458	
			present invention are one or more		
			polynucleotides comprising a		

1540 HCYBG67	10 876589	nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 923 of SEQ ID NO:1539, b is an integer of 15 to 937, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1539, and where b is greater than or equal to a + 14. Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 357 of SEQ ID NO:1540, b is an integer of 15 to 371, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1540, and where b is greater than or equal to a + 14. Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 892 of SEQ ID NO:1541, b is an integer of 15 to 906, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1541, and where b is greater than or equal to a + 14.	AA305259, L37080, Z47553 AA446378, AA305361, AA502360, A1912345, AA903395, AW377671, D80522, D81026, D80133, AW177440, AW360811, AW375405, A1262837, D80248, AW17511, AW352117, D80551, D80269, AW366296, D80366, D58283, D59859, D80022, C14331, D80166, D80366, D58283, D59859, D80022, C14331, D80166, D80360, D51799, D80391, D80164, D59275, D80240, D80253, D80043, D59787, D80227, D59502, AW375832, AW360844, D8130, AW360817, D80212, AW375832, AW37672, AW179023, AW378534, D80219, AA305578, C15076, D80038, D59610, D57483,
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		D51060, AW176467, D80378, AW352171, AW377676,
		AW179019, AW179024, AA514188, C14014, D80241,
		AW178906, AW352158, AW177505, AW179020,
		AW178909, AW177456, AA514186, AW179329,
		80, AW177733,
		AW178754, AW179018, D80132, AW178983, AW179004,
		75259, AW36
		, D8043
	-	_
		7723, AW35217
), AW378533, AW1789
		AI525913, AI525923,
		,681
		A67220, D897
		D26022, D88547,
		X67155, AF058696, A25909, Y12724, AR008278,
		6488, A82
		AB012117, AR016514, D50010, AR060138, A45456,
		I18367, A26615, AR052274, Y09669, AR060385,
		9, AR066487, AR038669, A431
		A30438, A85396, D88507, AR066482, A44171,
		A85477, I19525,
		3, X93549,
		U46128, AR008382
Pref	Preferably excluded from the	AA307067, AA827296, AA307068, AA972507,
pres	present invention are one or more	AA074169, AL134865, AA096156, AA247393,
poly	polynucleotides comprising a	AA091519, I81218, U30872, U19769, I35495,
nucl	nucleotide sequence described by	AF194970
the	the general formula of a-b, where a	
1.8	is any integer between 1 to 965 of	
SEQ	SEQ ID NO:1542, b is an integer of	
15	to 979, where both a and b	
corr	correspond to the positions of	

			de residues show	
			NO:1542, and where b is greater than or equal to a + 14.	
1543	HCYBI92	876592	Preferably excluded from the	R24666, AA305450, M63635, M64590, D90239
		_	present invention are one or more	
			polynucleotides comprising a	
			nucleotide sequence described by	
			the general formula of a-b, where a	
			is any integer between 1 to 287 of	
			SEQ ID NO:1543, b is an integer of	
			15 to 301, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1543, and where b is greater	
			than or equal to a + 14.	
1544	HWMCC2	876595	Preferably excluded from the	AI690065, AI480300, AA927896, AI288678,
	∞		present invention are one or more	AI343570, AI343569, AI678924, AW339479,
	•		polynucleotides comprising a	AA836387, AA836420, AC006011
			nucleotide sequence described by	
			the general formula of a-b, where a	
		-	is any integer between 1 to 638 of	
			SEO ID NO:1544, b is an integer of	
		dd.	15 to 652, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1544, and where b is greater	
	;		than or equal to a + 14.	
1545	HWMAN6	876596	Preferably excluded from the	_
			present invention are one or more	AA315469, AA568218, AI150400, AA583146,
			\vdash	AW374998, AI955582, AW374874, AI832775,
			nucleotide sequence described by	AA345780, AA295520, AW360893, AA294858,
			the general formula of a-b, where a	AI445680, AW360892, AW360931, AA295782,
			is any integer between 1 to 2222 of	AF102542, AF038650, R32988, H99036, N39174,
			SEQ ID NO:1545, b is an integer of	N45249, N62843, W60278, W79341, W79441, W93292,
			15 to 2236, where both a and b	W93293, W92077, W92073, AA083227, AA102315,
			correspond to the positions of	AA111889, AA121668, AA121740, AA505444,

MATCHE Match Mat				1	
HCQCR04 876597 Preferably excluded from the present invention are one or more plantage and where a is any integer between 1 to 342 of SEQ 18				de residues shown in SEQ	, AA574144, AA738177,
HCQCR04 876597 Preferably excluded from the polymuclacides comprising a nucleotide sequence described by the general formula of a -b, where a is any integer by correspond to the positions of nucleotide sequence described by the general formula of a -b, where a is any integer between 1 to 342 of SEQ 1D NO.1546, and where b is greater than or equal to a + 14. HWMF448 876600 Preferably excluded from the positions of nucleotide residues shown in SEQ 1D plymucleotide residues one or more plymucleotide sequence described by nucleotide sequence described by a 1302190, AIS9087. AA31099 AM88539. AA31099 AM88539 AM88530				and where b is	, AA/81330, AIU15034,
HCQCR04 876597 Preferably excluded from the present invention are one or more polynuclectides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between it to 142 of SEQ ID NO:1546, b is an integer of nucleotide residues shown in SEQ ID NO:1546, and where b is greater than or equal to a + 14. AA8613252, AI911238, AI186148, AI743777, AA861325, AI911238, AI186148, AI743777, AA861325, AI911238, AI186148, AI743777, AA861325, AI911238, AI186148, AI743777, AIRAGE SEGO ID NO:1546, and where b is greater than or equal to a + 14. AA861325, AI911106, AI092279, AA69822, AI886794, AI886784, AI886784, AI886784, AI886784, AI89784,				equal to a	, AI360138, AI383772,
HCQCR04 876597 Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a b, where a is any integer between 1 to 342 of SEQ ID NO.1546, b is an integer of 15 to 356, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO.1546, b is an integer of 15 to 356, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO.1546, and where b b is greater than or equal to a + 14. HWMFE48 876600 Preferably excluded from the 10 158 of 100.1540, A1380352, A1381523, A1381648, A138713, A138186148, A138713, A1381865, A1381865, A13818653, A13818653, A1381865, A13818653, A13818653, A13818653, A13818653, A13818653, A138186148, A138713, A1381865, A1381865, A13818653, A1381865, A1381865, A13818653, A1381863, A13818653, A13818					, AI127637, AI129439,
HCCCR04 876597 Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 342 of SEQ ID No:1546, b is an integer of nocleotide residues shown in SEQ ID No:1646, and where b is greater than or equal to a + 14. HWAMFE48 876600 Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide residues shown in SEQ ID No:1547, b is an integer of sequence described by nucleotide requence described by the general formula of a-b, where a nucleotide residues shown in SEQ ID No:1547, and where b is greater No:1547, and where b is greater HWTBN44 876601 Preferably excluded from the present invention are one or more No:1547, and where b is greater D1707, U17080 No:1547, and where b is greater No:1547, and where b is greater D1707, U17080 No:1547, and where b is greater No:1547, and where b is greater D1707, U17080 No:1547, and where b is greater No:1547, and where b is greater D1707, U17080 No:1547, and where b is greater D1707, U17080 No:1547, ANSESSER ANSES					, AI208460,
present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 342 of seq is any integer between 1 to 342 of seq is any integer between 1 to 342 of seq is and where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1546, and where bis greater than or equal to a + 14. HWMFE48 876600 Preferably excluded from the polynucleotide sequence described by nucleotide sequence described by any integer between 1 to 1158 of any any any a	1546	HCQCR04	876597	Preferably excluded from the	
polymucleotides comprising a nucleotide sequence described by the general formula of a -b, where a is any integer between 1 to 342 of SEQ ID No.1546, b is an integer of nucleotide residues shown in SEQ ID nucleotide residues shown in SEQ ID nucleotide comprising a present invention are one or more proportional of a -b, where a polymucleotide comprising a nucleotide residues shown in SEQ ID nucleotide residues shown in SEQ ID nucleotide residues comprising a nucleotide residues shown in SEQ ID		,		nvention	
the general formina of a-b, where a is any integer between 1 to 342 of SEQ ID No:1546, b is an integer of 15 to 356, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID No:156, and where b is greater than or equal to a + 14. HWMFE48 876600 the feetably excluded from the polymorleotides comprising a polymorleotides comprising a polymorleotide sequence described by the general formina of a-b, where a mucleotide sequence described by a pass 1730, A1302103, A1302179, the general formina of a-b, where a mucleotide residues shown in SEQ ID No:154, b is an integer of A1502104, A1720296, A1503096, A3030106, A503017, A1003179, and where both a and b correspond to the positions of nucleotide residues shown in SEQ ID No:1547, and where b is greater A4733240, AA513017, A10321701, than or equal to a + 14. HMTBN44 876601 Preferably excluded from the present invention are one or more polymorleotides comprising a present invention are one or more polymorleotides equence described by the qeneral formula of a-b, where a the qeneral formula of a-b, where a the deneral formula of a-b, where a the present invention are one or more polymorleotides equence described by the qeneral formula of a-b, where a the quality of a-b, where a the q				otides comprising	
the general formula of a-b, where a is any integer between 1 to 342 of SEQ ID No:1546, bis an integer of 15 to 356, where both a and b correspond to the positions of no:1546, and where b is greater than or equal to a + 14. HWMFE48 876600 Preferably excluded from the polyuucleotides comprising a nucleotide sequence described by present invention are one or more polyuucleotides comprising a nucleotide sequence described by the general formula of a-b, where a Na02130, A1951039, A191179, SEQ ID No:1547, b is an integer of A186718076, A552103, A1911706, A1911706, A1911707, A1911706, A1911707,				nucleotide sequence described by	
is any integer between 1 to 342 of SEQ ID NO:1546, b is an integer of IS to 356, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1546, and where b is greater than or equal to a + 14. HWMFE48 876600 Preferably excluded from the positions of present invention are one or more preferably excluded from the prostition of present invention are one or more described by an integer of a 1922396, A191105, A198037, A1981173, A198117				of a-b, where	
SEQ ID NO:1546, b is an integer of 15 to 356, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID (NO:1546, and where b is greater than or equal to a + 14. HWMFE48 876600 Preferably excluded from the positions of nucleotide sequence described by the general formula of a - b, where a nucleotide sequence described by the general formula of a - b, where a nucleotide residues shown in SEQ ID (NO:1547, and where both a and b correspond to the positions of nucleotide residues shown in SEQ ID (NO:1547, and where b is greater han or equal to a + 14. HWTBN44 876601 Preferably excluded from the positions of nucleotide sequence described by the general formula of a - b, where a nucleotide sequence described by the general formula of a - b, where a nucleotide sequence described by the general formula of a - b, where a nucleotide sequence described by the general formula of a - b, where a nucleotide sequence described by the general formula of a - b, where a nucleotide sequence described by the general formula of a - b, where a nucleotide sequence described by the general formula of a - b, where a nucleotide sequence described by the general formula of a - b, where a nucleotide sequence described by the general formula of a - b, where a nucleotide sequence described by the general formula of a - b, where a nucleotide sequence described by the general formula of a - b, where a nucleotide sequence described by the general formula of a - b, where a nucleotide sequence described by the general formula of a - b, where a nucleotide sequence or more nucleotide sequence described by the general formula of a - b, where a nucleotide sequence described by a nucleotide sequence or more nucleotide sequence described by a nucleotide sequence described by a nucleotide sequence or a - b, where a nucleotide sequence described by a nucleotide sequence or a - b, where a nucleotide sequence described by a nucleotide sequence or a - b, where a nucleotide sequence described by a nucleotide sequence or a - b, w				342	
15 to 356, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1546, and where b is greater than or equal to a + 14. HWMFE48 876600 Preferably excluded from the positions of a polynucleotide sequence described by the general formula of a-b, where a polynucleotide sequence described by the general formula of a-b, where a A1302130, A1255082, AA158037, A1991179, is any integer between 1 to 1158 of A4594818, A167941, A1738706, AA544173, SEQ ID NO:1547, b is an integer of A4887784, A4552303, A1424977, A1020179, NO:1547, and where b is greater than or equal to a + 14. HMTBN44 876601 Preferably excluded from the polynucleotides comprising a A1446030, D62937, AA15177, AA15089, AA151869, Dolynucleotide sequence described by the general formula of a-b, where a nucleotide sequence described by the general formula of a-b, where a nucleotide sequence described by the general formula of a-b, where a nucleotide sequence described by the general formula of a-b, where a					
nucleotide residues shown in SEQ ID NO:1546, and where b is greater than or equal to a + 14. HWMFB48 876600 Preferably excluded from the population are one or more described by the general formula of a-b, where a his any integer between 1 to 1158 of a passible and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1547, and where b is greater than or equal to a + 14. HMTBN44 876601 Preferably excluded from the polymorbotide sequence described by a passible and b and b and and and and b and			_	15 to 356, where both a and b	
No.1546, and where b is greater				correspond to the positions of	
HWMFE48 876600 Preferably excluded from the present invention are one or more polynucleotides comprising a polynucleotide sequence described by the general formula of a-b, where a polynucleotide sequence described by the general formula of a-b, where a polynucleotide sequence described by the general formula of a-b, where a polynucleotide sequence described by the general formula of a-b, where a polynucleotide residues shown in SEQ ID NO:1547, b is an integer of a pay 175, b				shown in SEQ	
HWMFE48 876600 Preferably excluded from the present invention are one or more polynucleotides comprising a polynucleotide sequence described by a present invention are one or more properly and a polynucleotide sequence described by the queryal formula of a-b, where both a and b				is	
HWMFE48 876600 Preferably excluded from the present invention are one or more pleases				equal to a + 14.	
present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a sequence described by the general formula of a-b, where a sequence described by the general formula of a-b, where a sequence described by the general formula of a-b, where a sequence described by the greater invention are one or more polynucleotide sequence described by the general formula of a-b, where a polynucleotide sequence described by the greater invention are one or more polynucleotide sequence described by the greater invention are one or more polynucleotide sequence described by the greater invention are one or more polynucleotide sequence described by the greater invention are one or more polynucleotide sequence described by the greater invention are one or more polynucleotide sequence described by the greater invention are one or more polynucleotide sequence described by the general formula of a-b, where a	1547	HWMFE48	876600	y excluded from	2, AI911238, AI186148,
polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a nucleotide sequence described by the general formula of a-b, where a has any integer between 1 to 1158 of A887784, A158092, A4158037, A1991179, SEQ ID NO:1547, b is an integer of correspond to the positions of nucleotide residues shown in SEQ ID A78392, D31212, T65680, A465630, AA15804, AA15809, AA382098, NB NO:1547, and where b is greater than or equal to a + 14. HMTBN44 876601 Preferably excluded from the polynucleotide sequence described by the general formula of a-b, where a				present invention are one or more	0, AI004989, AI808771, AA83855
MATBN4 876601 Preferably excluded from the polynucleotide sequence described by the general formula of a-b, where a his any integer between 1 to 1158 of 1954918, A1167941, A1738706, AA524173, SEQ ID NO:1547, b is an integer of correspond to the positions of nucleotide residues shown in SEQ ID NO:1547, and where b is greater than or equal to a + 14. HMTBN4 876601 Preferably excluded from the polynucleotide sequence described by the general formula of a-b, where a process and the polynucleotide sequence described by the general formula of a-b, where a process and the process and the polynucleotide sequence described by the general formula of a-b, where a process and the process and th				polynucleotides comprising a	5, AI911106, AI092279,
the general formula of a-b, where a highest petween 1 to 1158 of SEQ 10 NO:1547, b is an integer of correspond to the positions of nucleotide residues shown in SEQ ID NO:1547, and where b is greater than or equal to a + 14. HMTBN44 876601 Preferably excluded from the polynucleotide sequence described by the general formula of a-b, where a high sand in sequence described by the polynucleotide sequence described by the sequence of the sequence of the sequence of the sequence described by the sequence of the sequence of the sequence o				nucleotide sequence described by	6, AI955005, AI034008,
is any integer between 1 to 1158 of AA887784, AA552303, AI424977, AI024177, SEQ ID NO:1547, b is an integer of AA887784, AA552303, AI424977, AI024177, AI02117, AI02117, AI021107, Were both a and b correspond to the positions of nucleotide residues shown in SEQ ID AA778392, D31212, T65680, AA45509, AA151201, than or equal to a + 14. HMTBN44 876601 Preferably excluded from the polynucleotides comprising a nucleotide sequence described by the equeral formula of a-b, where a					, AI285082, AA158037,
SEQ ID NO:1547, bis an integer of AA857384, AA552303, AI424977, AI024177, AI024177, AI021807, William or the positions of nucleotide residues shown in SEQ ID AA78392, D31212, T65680, AA465630, AA15 NO:1547, and where b is greater than or equal to a + 14.				is any integer between 1 to 1158 Of	8 AT167941 AI738706.
15 to 1172, where both a and b AS594882, W85752, AA315096, AI672956, R9 correspond to the positions of nucleotide residues shown in SEQ ID AA778392, D31212, T65680, AA465630, AA15 N0:1547, and where b is greater than or equal to a + 14. HMTBN44 876601 Preferably excluded from the polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a				coger permitted to the contract of the contrac	4 AASS303 AT424977
Correspond to the positions of nucleotide residues shown in SEQ ID AA778392, D31212, T65680, AA465630, AA15 NO:1547, and where b is greater than or equal to a + 14. HMTBN44 876601 Preferably excluded from the polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a				יייטייייי לייסיליייייייייייייייייייייייי	WEEZAL ATTOORS AT
Correspond to the positions of nucleotide residues shown in SEQ ID AA778392, D31212, T65680, AA465630, AA15 NO:1547, and where b is greater than or equal to a + 14. HMTBN44 876601 Preferably excluded from the polynucleotides comprising a nucleotide sequence described by the qeneral formula of a-b, where a				ובס רח בדין אוופדב מסרוו מ מווח מ	COCCOCING COCCICCES CUCUCIN
nucleotide residues shown in SEQ ID AA778392, D31212, T65680, A4465630, AA15 NO:1547, and where b is greater than or equal to a + 14. HMTBN44 876601 Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the qeneral formula of a-b, where a					, 20108W ,
NO:1547, and where b is greater AA641295, AA928364, AA812254, AI351201, than or equal to a + 14.			_	e residues shown in SEQ	, D31212, T65680, AA465630, AALS
than or equal to a + 14. AW382084, AI383689, AA215354, AI873941, AW382340, AA639464, AW382339, AW351859, U17079, U17080 Present invention are one or more polynucleotides comprising a nucleotide sequence described by the qeneral formula of a-b, where a				and where b is	, AA928364, AA812254, AI351201
HMTBN44 876601 Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the qeneral formula of a-b, where a				or equal to a + 14.	AI383689, AA215354, AI873941,
HMTBN44 876601 Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a					AA639464, AW382339,
HMTBN44 876601 Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the qeneral formula of a-b, where a		_			U17079, U17080
present invention are one or more D79906, AW151367, polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a	1548	HMTBN44	876601	Preferably excluded from the	D62937, AA344217,
otides comprising a e sequence described by al formula of a-b, where	•			nvention are one or	AW151367,
described by of a-b, where				polynucleotides comprising a	
of a-b, where				nucleotide sequence described by	
				of a-b, where	

		10:1548, b is an i	
		15 to 1423, where both a and b	
		correspond to the positions of	
		nucleotide residues shown in SEQ ID	
		NO:1548, and where b is greater	
		equal to a + 14.	
1549 HCRO104	104 876602	Preferably excluded from the	M63806, AF035406, M96066, S68616
		present invention are one or more	
		polynucleotides comprising a	
		nucleotide sequence described by	
		the general formula of a-b, where a	
		is any integer between 1 to 443 of	
	_	correspond to the positions of	
_		nucleotide residues shown in SEQ ID	
		NO:1549, and where b is greater	
		equal to a + 14.	: : : : : : : : : : : : : : : : : : : :
1550 HTWCT64	T64 876608	Preferably excluded from the	AW118825, AI582268, AI924840, AI686918,
		present invention are one or more	AI689468, AI565967, AI471821, AW167093,
	_	polynucleotides comprising a	AW438815, AI560103, AW192267, AI351758,
_		nucleotide sequence described by	AI204255, AA948069, AA775662, AI160736,
		the general formula of a-b, where a	AA975121, AI347454, AW381442, AI086345,
		yer between	AI805695, AA441899, AW132052, AA233648,
_			AW204634, AI470694, AA464178, AA693693,
			AI061108, AW028857, N90723, AI275105, AI290106,
		correspond to the positions of	AW130518, N33172, AA031928, AA476308, AI682854,
		nucleotide residues shown in SEQ ID	AI358603, AI332311, AW381443, AI696369,
		NO:1550, and where b is greater	AW381398, AI472619, AI383588, AA404636,
		equal to a + 14.	AA180763, AA233637, AW381420, AA032029,
			AI559765, N90350, N44956, W06927, AA182891,
	-		C05190, AA883620, AI696426, AA618268, D90034,
			E01793, E01792, E01791, D28915, D28914, D28912
1551 HETBI79	179 876609	Preferably excluded from the	AI346674, AI348020, AI890197, AW291166,
		present invention are one	AA700159,

			polynucleotides comprising a	AA535792,	N76634, AA815232, A	AI343929, AA490536,
			nucleotide sequence described by			
			the general formula of a-b, where a	AA809480,	AI318395, AI761658,	AI140011,
			is any integer between 1 to 2526 of	AW190983,	AA488989	AW291783,
			SEQ ID NO:1551, b is an integer of	AI285896,	AA627444, R84232, A	AI674736, AI280867,
			15 to 2540, where both a and b	H72489, AA	H72489, AA488770, AA813879, A	AI685538, AI858181,
			correspond to the positions of		AA167381, N54554, N	N54554, N71216, AA971023,
			nucleotide residues shown in SEQ ID	AA704201,	AI612846, AW294335, N22015,	N22015, R10105,
			NO:1551, and where b is greater	AA744665,	AI680111, AI361708, AA313609, N755	AA313609, N75553,
			a)	AA337910,	H72889, AI689838, R87634,	187634, AI867541,
				AW015119,	R38671, R00317, AAS	R00317, AA548940, AI886417,
		-		T98789, WO	T98789, W05347, AA337673, T98	T98788, F10720,
				- 1	AW374/6/, AC004687	
1552	HWTBM65	876610	Preferably excluded from the		AI686316, AW137243, AW193522	AW193522,
			present invention are one or more	AW373055,	D79340, AI796896, AC004079	C004079
			polynucleotides comprising a			
			nucleotide sequence described by			
			is any integer between 1 to 594 of			
			SEQ ID NO:1552, b is an integer of			
			15 to 608, where both a and b	-		
			correspond to the positions of			
			nucleotide residues shown in SEQ ID			
			NO:1552, and where b is greater			
			than or equal to a + 14.			
1553	HCQBN77	876612	Preferably excluded from the	AA908796,	AA431249, AI743453,	A1433466,
			present invention are one or more		AW302156, AA758918,	AA595771,
			polynucleotides comprising a	AA432263,	AA887241, AI459626,	AA931083,
			nucleotide sequence described by	AI522039,	AA707461, AI612992,	AA834959, R50375,
			the general formula of a-b, where a	AI004115,	AI203186, R48003, R	R48117, L47334,
			teger between 1 to 770	AC005324,	AA976609	
			SEQ ID NO:1553, b is an integer of			
			15 to 784, where both a and b			
			correspond to the positions of			
			nucleotide residues shown in SEQ ID			
			NO:1553, and where b is greater			

			than or equal to a + 14.	
1554	HKAED74	876621	ly excluded from invention are one cotides comprising ectides comprising all formula of anteger between 10:1554, b is an integer between and to the positic de residues shown and where b is gequal to a + 14.	0, AA478680, AI972505, A 8, AI923250, AA210747, A 2, AA418404, AI683375, A 7, AA668890, AA315808, A 5, AI469242, AA814749, A 1, AA171797, AI745538, A 1, AI831534, AI206300, A 3, W95561, AI189412, AA6 0, AI199241, T75325, AI0 5, AA406142, F12995, F13 AA424821, T90046, T1928 8, F10590, AI868932, AA2 AA367325, AA428537, AA2 1, AI793116, AI793143, R
1555	нс <u>о</u> дт20	876622	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 380 of SEQ ID NO:1555, b is an integer of 15 to 394, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:155, and where b is greater than or equal to a + 14.	D60051, H57196, AI125536
1556	HCRMD40	876630	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 332 of SEQ ID NO:1556, b is an integer of 15 to 346, where both a and b	AL044257, W40373, AW250560, AA643353, AI991172 AA402608, AW249124, AI554578, AW328561, AW246456, AW051430, AA308337, AI346750, AW166193, AA703840, AI143755, AI951822, AW080812, AI189652, AI885695, AW166148, AW082817, AI953814, AA602780, AI951334, AI191618, AW248692, W45258, AA503856, AI378866 AA916922, AI089026, AA599791, AA032143, H48844

_			correspond to the positions of	AA402390 AT192449 AA826583	826583 AW	OFFER 7CAOTOMA
			s shown ir			
	_		and where b is greater			
			equal to a + 14.			
1557	HFIHO78	876631	Preferably excluded from the	AW150197, AA846471, AI	AI146351, AI	AI276560, H96798,
			present invention are one or more	AW016664, AA253395, W07219, H97716, M63896	7219, H977	16, M63896,
			polynucleotides comprising a	L13853, S74227, L06865		
			nucleotide sequence described by			
			the general formula of a-b, where a			
			is any integer between 1 to 1563 of			
·			SEQ ID NO:1557, b is an integer of			
			15 to 1577, where both a and b			
			correspond to the positions of			
			nucleotide residues shown in SEQ ID			
			NO:1557, and where b is greater			
:			than or equal to a + 14.			
1558 F	HCRPG35	876633	Preferably excluded from the	AC004030		
_			present invention are one or more			
			polynucleotides comprising a			
_			nucleotide sequence described by			
			the general formula of a-b, where a			
			is any integer between 1 to 264 of			
			SEQ ID NO:1558, b is an integer of			
			15 to 278, where both a and b			
			correspond to the positions of			
			nucleotide residues shown in SEQ ID			
			NO:1558, and where b is greater			
			than or equal to a + 14.			
1559 F	HSQFQ92	876637	Preferably excluded from the	, AI692181,	A1275606, AI	AI453065,
			present invention are one or more	AI521837, AI634107, AW	AW130839, AI	AI654841,
			polynucleotides comprising a	AA424967, AA059190, AA	AA047896, AA	AA148675,
			nucleotide sequence described by	AW085538, AA026771, AI	AI261336, AI	AI696507,
			the general formula of a-b, where a	AA992863, N66291, R85666	99.	
			is any integer between 1 to 737 of			
			ın inte			
			15 to 751, where both a and b			

			correspond to the positions of				
			nucleotide residues shown in SEQ ID				
			NO:1559, and where b is greater				
			than or equal to a + 14.				
1560	HUFBF32	876638	Preferably excluded from the	AL134555,	AI925308, A	AI625207,	AI969783,
			present invention are one or more	AW262828,	2	AI685887,	AA206222,
			polynucleotides comprising a	AI086025,	_	AA143639,	AI268485,
			nucleotide sequence described by	AI312871,	AL134554, A	AA969162,	AI282923,
			the general formula of a-b, where a	AA074267,	AA206652, N	33991, N2	N33991, N22039, T09372,
			is any integer between 1 to 1924 of	AI760417,	AA146631, A	W083343,	AW083343, AI479411,
			SEQ ID NO:1560, b is an integer of	AA742178,	AW054790, A	AI586977, AI948545	AI948545,
				AI991591,	T59451, AI565918,	65918, AI	AI572624, AA627495,
			correspond to the positions of	AA236672,	AI798559, AW291470, AA29244	W291470,	AA292449,
			nucleotide residues shown in SEQ ID	AA593202,	T58112, AI8	15717, AI	T58112, AI815717, AI698280, AI432649
			NO:1560, and where b is greater				
			than or equal to a + 14.				
1561	HTXC005	876643		AW411282,	R08081, AA3	AA307047, T9	T98713, AW351792,
			present invention are one or more	AA325934,	AW375839, AI694682, AI968390,	1694682,	AI968390,
			polynucleotides comprising a	AW370749,	AW370756, U	U43431	
			nucleotide sequence described by				
			the general formula of a-b, where a				
			is any integer between 1 to 875 of				
			SEQ ID NO:1561, b is an integer of				
			15 to 889, where both a and b				
			correspond to the positions of				
			nucleotide residues shown in SEQ ID				
			NO:1561, and where b is greater				
			than or equal to a + 14.				
1562	HWMBJ09	876645	Preferably excluded from the	AW337919,	AA523430, A	AL044577,	AW194215,
			present invention are one or more	AI686556,	AI671043, A	AA652193,	AI815222,
			polynucleotides comprising a	AI694846,	AA480192, A	AI289064,	AI910616,
			nucleotide sequence described by	AI923986,	AI557645, A	AI799943,	AI077441,
	-		the general formula of a-b, where a	AW007863,		AI123788,	AW024224,
			is any integer between 1 to 1371 of	AI355044,	AW130857, A	AW054917,	AA552445,
			SEQ ID NO:1562, b is an integer of	AA923164,	AA300093, A	AI686879,	AI240984,
			15 to 1385, where both a and b	AI625429,	AI446337, A	AI557649,	AI557647,

			correspond to the positions of nucleotide residues shown in SEQ ID	AA524488, AI557652, AI557651, AI557653, AA579950, AW338240, AI557650, AA480098,
			NO:1562, and where b is greater	6, AIS57654, AIS57655, AIS57648,
_			than or equal to a + 14.	AA994813, AL044578, AI383197, AA910275, K05862, AA887744. R05776. AI940377. AA594829. AA858443.
				AW337931, AW057864, AI720420,
				AI557646, AW363060, X87342
1563	HSIDP84	876646	Preferably excluded from the	W316845, AI
			present invention are one or more	AI623768, AI934315, AI692242, AI023791,
			polynucleotides comprising a	A1935868, A1934327, AI818628, AI589269,
			nucleotide sequence described by	C05899,
_			the general formula of a-b, where a	AI703259, H70829, AI598076, H61582, H70828,
			is any integer between 1 to 848 of	AI932542, AI582914, AI587377, AI565896,
			SEQ ID NO:1563, b is an integer of	AI445979, H94487, H79481, AI888892, H61583,
			15 to 862, where both a and b	M84424, J05036
			correspond to the positions of	
•			nucleotide residues shown in SEQ ID	
			NO:1563, and where b is greater	
			than or equal to a + 14.	
1564	HUSJA29	876647	Preferably excluded from the	_
			present invention are one or more	
			polynucleotides comprising a	AA694059,
			nucleotide sequence described by	
			the general formula of a-b, where a	
			is any integer between 1 to 3093 of	
			SEQ ID NO:1564, b is an integer of	
			15 to 3107, where both a and b	AI418973, N94584, N22975, AW009450, AI423399,
_			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1564, and where b is greater	AI399686, W49640, AI280345, AA703127, AI632111,
			equal to a + 14.	T63353, AI865130, AI474045, H47786, AI274468,
				AI341413, AW016684, AI399864, AA694012,
				AI097106, AL040613, AW182238, AA431110, R14723,
				Ø
				, W23791,
				AA972808, Z45677, R36481, AA479212, AI567031,

				R62535, R84588, N50507, AA969851, T97034, AA649044, AA315207, AA649043, AI471105,
				AIO86675, R36482, AA613263, AIO51650, Z41345, B42442 AIO74320 R66089 AA812544 R06604
				T96927, R06660, N32390, AI868697, R06669,
				AA432124, N79367, T63677, Z20112, AA883725,
				AI220180, AC004711, AB020684, AJ011911, AC005271, A74567, AA770028
1565	HCOAG09	876648	Preferably excluded from the	AF084644,
<u> </u>			present invention are one or more	AF188476, AF182217, AJ009937
			polynucleotides comprising a	
			nucleotide sequence described by	
-			the general formula of a-b, where a	
				-
		_	SEQ ID NO:1565, b is an integer of	
•		_	15 to 300, where both a and b	
	-	_	correspond to the positions of	
-			nucleotide residues shown in SEQ ID	
	-		NO:1565, and where b is greater	
			than or equal to a + 14.	
1566	HCROT53	876649	Preferably excluded from the	U17105, Z36714, U20612, Z47766, U20636
			present invention are one or more	
			otides comp	
	-		nucleotide sequence described by	
	•		wher	
			:1566, b is an inte	
			15 to 537, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1566, and where b is greater	
			than or equal to a + 14.	
1567	HOENX50	876652	Preferably excluded from the	AF039023, AC006432
			present invention are one or more	
			polynucleotides comprising a	
			nucleotide sequence described by	

		the general formula of a-b, where a	
		SEQ ID NO:1567, b is an integer of	
		15 to 333, where both a and b	
		correspond to the positions of	
		nucleotide residues shown in SEQ ID	
		NO:1567, and where b is greater	-
		than or equal to a + 14.	
1568 HCEOW20	720 876656	Preferably excluded from the	AA985339, AA325781, AA041430, AC005531
		present invention are one or more	
		polynucleotides comprising a	
		nucleotide sequence described by	
	-	the general formula of a-b, where a	
		SEQ ID NO:1568, b is an integer of	
		15 to 649, where both a and b	
		correspond to the positions of	
		nucleotide residues shown in SEQ ID	
		NO:1568, and where b is greater	
		than or equal to a + 14.	
1569 HCRMG16	116 876657	Preferab	299757
		present invention are one or more	
	_	polynucleotides comprising a	
		nucleotide sequence described by	
	•	the general formula of a-b, where a	
		is any integer between 1 to 379 of	
		SEQ ID NO:1569, b is an integer of	
		15 to 393, where both a and b	
		correspond to the positions of	
	-	nucleotide residues shown in SEQ ID	
		NO:1569, and where b is greater	
		than or equal to a + 14.	
1570 HCEPH79	79 876660	Preferab	AA326212
		eotides comprising a	
		nucleotide sequence described by	

			the general formula of a.h where a	
			general retinate of a straint integer between 1 to	
			o:1570. b is an inte	
			6, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1570, and where b is greater	
1571	HFOYY56	876666	Preferably excluded from the	AI828664, AW189077, AA186731, AA058868,
			present invention are one or more	AA723578, AL121358, AI221227, AI093392,
			polynucleotides comprising a	AI138553, AW019870, AI803661, AA826404,
			nucleotide sequence described by	AI004869, N67735, AI188839, AI474328, N64380,
•			the general formula of a-b, where a	T71617, AI630399, AL120719, AA127002, AW386045,
			is any integer between 1 to 1643 of	AA243169, N70412, N40572, AA977240, AI798975,
			SEQ ID NO:1571, b is an integer of	H41757, H41758, AL046756, H40420, H50495,
			15 to 1657, where both a and b	T91967, N44609, AA125926, H14602, AI950747,
			correspond to the positions of	H20721, H72253, R10731, AW382088, AA069491,
			nucleotide residues shown in SEQ ID	R44126, AI472460, AA045529, AA731653, AW366585,
			NO:1571, and where b is greater	AI148840, AI373402, W58735, N35135, AI889177,
_			than or equal to a + 14.	AA127021, H71690, AA069453, AA125758, AI312614,
				AB006965, AF000430, AF061795, AF151685,
	٠		-	
				AF020211, AF020213, AF132939
1572	HSXDG80	876668	Preferably excluded from the	N76733, H97908, AI765923, AA100164, AI161123,
			present invention are one or more	AI269285, N45309, AI379293, AA026656, AA425856,
			polynucleotides comprising a	H06713, AA628959, N54759, AA323052, AI123671,
			nucleotide sequence described by	R78485, AA317233, N88108, T92033, T84742,
			the general formula of a-b, where a	AW263910, AI400524, AA628884, AW275553,
			is any integer between 1 to 1172 of	AI039362, R78527, AA249635, AI041425, N52791,
				AI699248, AA223953, AI191006, N59264, AB020715
			15 to 1186, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1572, and where b is greater	
			than or equal to a + 14.	
1573	HHEUK77	876675	Preferably excluded from the	AA313261, AA300475, AA133237, AI768979,

			present invention are one or more	AA580098, AA233499, AA314374, AW408727,
			leotides comprising	AA094260, AI751632
			nucleotide sequence described by	
			the general formula of a-b, where a	
		_	is any integer between 1 to 711 of	
		_	SEQ ID NO:1573, b is an integer of	
			15 to 725, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1573, and where b is greater	-
			than or equal to a + 14.	
1574	HHEDO14	876677		, AL037493,
			present invention are one or more	
			ř	٦,
			nucleotide sequence described by	AA937974, AA634429, AI004727, AI299652,
			the general formula of a-b, where a	AA032043, AA862157, AI291351, AA862156,
			is any integer between 1 to 1121 of	AA181981, AA993666, AA991222, N52079, AA496026,
			SEO ID NO:1574, b is an integer of	AI000697, AI581889, AW342034, AI972961,
			15 to 1135, where both a and b	AA948363, AA258118, AI971556, N89925, AA041553,
			correspond to the positions of	H49505, AI017756, AA031961, W19241, F02366,
			nucleotide residues shown in SEQ ID	\sim
			NO:1574, and where b is greater	AA748836, AI262706, AA436938, AA877698,
			equal to a + 14.	AA187708, AA081668, H94003, H49504, H73988,
				AA244456, AA259104, H95020, AA082449, F11149,
				Y13120, U11822, X74145, X83579, X57239, X65070
1575	HKIMC75	876680	Preferably excluded from the	AA193161, T10237, H11797, D44986, R25550,
		1	present invention are one or more	T77684, R91095, H15636, Z42961, R17883,
			polynucleotides comprising a	AA371122, AL035427, AF035288, AC007262
			nucleotide sequence described by	
			the general formula of a-b, where a	
			is any integer between 1 to 845 of	
			SEQ ID NO:1575, b is an integer of	
			15 to 859, where both a and b	
			nucleotide residues shown in SEQ ID	

			NO:1575, and where b is greater					
			than or equal to a + 14.					
1576	HWMB136	876683	Preferably excluded from the	AI435038,	Ċ	AI701595,	AI628945,	
			present invention are one or more	AI819240,	AI361891, A	AI057030,	AI808292,	
			polynucleotides comprising a	AI478205,	AA933801, 1	AA633552,	AI830350,	
			nucleotide sequence described by	AA513475,	AI093856, 1	AI566604,	AI559922,	
			the general formula of a-b, where a	AI000612,	AA587035, 1	AI222881,	T27670, AI308944	4,
			is any integer between 1 to 718 of	AI308779,	AA948404, 1	AI346156,	AA857101,	
			SEQ ID NO:1576, b is an integer of	AI539010,	AI871676, 1	AI628889,	AI344797,	
			15 to 732, where both a and b	AA865820,	AI658897, 1	AI475182,	AW082952,	
			correspond to the positions of	AW102783,	AI346307, 1	AI972243,	AL045929,	
			nucleotide residues shown in SEQ ID	AI682106,	AI344182, 1	AI590482,	AI345860,	
			NO:1576, and where b is greater	AI569870,	M16937, S4	S49765		
			than or equal to a + 14.					
1577	HE8TM64	876685	Preferably excluded from the	AI751497,	W25812, AA	AA307338, AA	AA305326, AI367808	- 8,
			present invention are one or more	AA332338,	AA545813, AA047778, AI25178	AA047778,	AI251787,	-
			Ч	AL045193,	D30819, AA319757, AW293922,	319757, AV	1293922, X68199,	
			nucleotide sequence described by	X69987, L	X69987, L00923, AJ001381, AJ001382	1381, AJOC	1382	
			the general formula of a-b, where a					
			is any integer between 1 to 1622 of					
			SEQ ID NO:1577, b is an integer of	-				
			15 to 1636, where both a and b					_
			correspond to the positions of					
			nucleotide residues shown in SEQ ID					
			NO:1577, and where b is greater					
			than or equal to a + 14.				-	
1578	HKLSA57	876687	Preferably excluded from the					
			present invention are one or more					
			polynucleotides comprising a					
			nucleotide sequence described by					
			the general formula of a-b, where a					
			is any integer between 1 to 645 of					
			SEQ ID NO:1578, b is an integer of					
			15 to 659, where both a and b			•		
			correspond to the positions of					
			nucleotide residues shown in SEQ ID					

			NO:1578, and where b is greater than or equal to a + 14.	
1579	HOGCV45	876689	rab	1, AA316125, AA779730, AI34
			present invention are one or more	D82400, AI928195, R59543, R51409,
			polynucleotides comprising a	F11900, T65476, AA081963,
			nucleotide sequence described by	T65486, D82182, AA188083, X84373, AR031997
			al formula of a-b, where	
			is any integer between 1 to 1852 of	
			SEQ ID NO:1579, b is an integer of	
			15 to 1866, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1579, and where b is greater	
			Ψ.	
1580	HADCX04	876690	Preferably excluded from the	AI824012, AA768896, AI400750, AW291960,
			present invention are one or more	AA449520, AI446344, AI911295, AA482984,
			polynucleotides comprising a	AA677454, C75000, AA211913, AA449089, AL039130,
			nucleotide sequence described by	AI086104, AA809866, AA814760, AA206769, R51297,
			the general formula of a-b, where a	Z40045, R59544, T65401, AW440101, AW197032,
			is any integer between 1 to 1482 of	AA280932, T65412, D81782, R59543, AI916155,
			SEQ ID NO:1580, b is an integer of	F09547, AA206804, AA304478, AA743706, C75037,
				AA209222, Z43988, R51409, F11900, AA316125,
			correspond to the positions of	T65476, X84373, AF053062
			nucleotide residues shown in SEO ID	
			NO:1580, and where b is greater	
			than or equal to a + 14.	
1881	HCRPH70	876693	Preferably excluded from the	AI452523, AI478635, AI744981, AI560901,
			present invention are one or more	
			polynucleotides comprising a	AA421151, AI660891, AW444552, AL039553,
		••	nucleotide sequence described by	AI745043, AI570244, AI333562, AA205872,
			the general formula of a-b, where a	AI719554, AI149680, AW439417, AI921227,
			is any integer between 1 to 3884 of	AA694055, AI601268, AA316992, AI393735,
			SEQ ID NO:1581, b is an integer of	AW190924, AA838650, AI269927, AI095118,
				AW151035, AI769469, AW337209, AI025693,
			correspond to the positions of	, AL039554,
			nucleotide residues shown in SEQ ID	AA936325, AI242821, AA814514, AL121252,

			NO:1581, and where b is greater	AW376485, AW131188, AW192413, AL121316,
			equal to a + 14.	AW014973, AA101068, AL039574, AW131134,
				6
	-		•	
				, AI537228, AA226093
			•	AA225947, AA397942,
			,	H25331, AA814957,
				AW380100, N75624, AA372640,
				AL046083, T54750,
				, AA352818, AI3077
				A344845, N22383, AA353560,
				AI762329, F01918, AA373973, T54663, N88370,
				AA206054, AI040829
			-	AI749924,
				T19805, AA082735, AW273597, AW374506, AI557427,
				AA857322, AI721273, AI423660, AA302091,
	•			AA181082, R17993, AW360799, H13417, AA977862,
				H13460, H13520, AW360925, AI206966, AI206949,
				9, X53586,
				X69902, X56559, AF166341, S66213, S66196,
				I32962, I32961, S52135, AF166343, AF166342
1582	HCRQM22	876696	Preferably excluded from the	AW403014, AI904490, AI831848, AA115313,
	,		present invention are one or more	AI761315, L16783, U74613, U83113, AR030545,
			polynucleotides comprising a	A79030, U74612, AC005841
			nucleotide sequence described by	
			the general formula of a-b, where a	
			ween 1 to 433	
			an inte	
			15 to 447, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1582, and where b is greater	
			than or equal to a + 14.	
1583	HKAEBIS	876697	Preferably excluded from the	AL036025, AW170264, AI752535, AI005255,
			present invention are one or more	AI983435, AW246157, AA830412, AA100899,

	polynucleotides comprising a	AW029286, AW249623, AI817149, AI188189,
	sedne	AI080559, AI351548, AI800612, AA053203,
	l formula	I472277, AA514834, AI805161, AW19053
	is any integer between 1 to 1260 of	AI674923, AI126935, AI692174, AW338703,
	NO:	90
	o 1274, where	AI280045, AA775722, AA748994, AW340009,
	correspond to the positions of	AW021825, AW079812, AA687655, AA157990,
	residue	AI335523, H28772, AA053118, AA179129, R98683,
	NO:1583, and where b is greater	F37299, AA490300, AA128782, AI222643, AI971507,
	equal to a + 14.	AA158221, W22913, AI808088, AI241313, AA128683,
		W75952, AA490392, AA937369, W70210, F27137,
		, R98910, AA878476, AA835695
		I698637, AA568407,
		AI873390, AA191377, AA352963, AA845387,
		T99184,
		AA190767, H19574, H92872, AA317262, H46433,
		AI356724,
		, AI864085, AA732079, AI701200, F312
		AW087408, T99183, AI345010, AW152550, AI890507,
), AA715307, AA809
		AI520946, AA761557, AI445992, AI659795,
		AA641818, AW075608, AA857847, AW327325,
		AI860674, AA748353, AW090087, AI567971,
		AI433976, AL045413, AI860783, AI963172,
		AI590043, AI624543, AI064830, AI440238, N29277,
		AL038529, AW088037, AL038645, AW075084,
		AI310925, AW161202, AI538885, AI828574,
		AW161579, AI567582, AI289791, AI471429,
		AL120700, AW151136, AA659314, AI539771,
		AL121270, AI432644, AW162194, AI537677,
		00659,
-		540674, AI815232, AI801325,
		0523, AI537617, AI538850, A
		AI270350, AI582932, AL043168, AI923989,

	AI872423, AI284517, AI500706, AI890576,
	237, AI491776, AW15113
	σ
	, AI582912, AI284509,
	66573,
-	14, AI866469,
	AI805769, AI434242, AI888661, AI312364,
	AIS00714, AI284513, AI345180, AI888118,
	5429,
	AI355779, AI889147, AI623736, AI581033,
-	AI371228, AI334884, AI491710, AI440252,
	AI431307, AW269098, AL047422, AW268251,
	AI114703, AI866786, AI860003, AI610557,
-	AI431316, AI433037, AI242736, AA808175,
	AI887499, AW151979, AI539781, AI364788,
	AI867068, AW268768, AI702065, AI539707,
	AI885949, AW089557, AI559957, AI285419,
	AW079432, AW089562, AI567953, AI815150,
	AI445620, AI671642, AI816055, AC004922, U26541,
	19367, U65960, U726
	I48978, AF132676, AF061836, AJ242859, Z72491,
	I89947, AF153205,
	AL122049, A08913, E02914,
	, AF017152, AL133049, AL110280,
	5, A18777, A77033, A77035, X70
	, AF061573, AL1330
	X93495, AF067790, I89931,
	D89079, A08911, AR038854,
	F113694, S83456, A07
	382, AF126488, AL023657, AL137533, X99
	AF102578, AL133619, X65873, E03671, AF079763,

10, I89934, AL117432, AL133565, AL133565, AL13361, AL137539, AL127110, AF100931, AL137555, AL133080, AL133081, AF192555, AT13753, AL133080, AL122121, A08911, A75801, AL137621, AL080163, AL122121, A08912, AL137621, AL133014, AL133014, AL133067, AL133077, AF118094, A65341, U58996, AL0800777, AF118094, A65341, U58996, AL133077, AL133057, AL133057, AL133057, AL133057, AL133057, AL133054, X6337, AL133053, AL133054, X6337, AL133053, AL133558, AL133054, X6337, AL133053, AL133558, AL133054, X6337, AL133558, AL133559, AL133558, AL133559, AL13359, AL133559, AL133559, AL133559, AL133559, AL133559, AL133559, AL	
s, ALU49300, A86538, AB029065, AF09799 3, Y11254, AR029490, AL122106, AF11185	
86558, AB029065, AF09799	_
985, I80064, AF114818, AL049464,	
901, AF137367, U35846, AC003032, ALL37	
471, AL049452, AF044323, AJ010277,	
528, AF118092, AF120268,	
5, AF106945, AF091084, X82434, X6686	
118, AL050092, X72387, AL050138, U42766	
AL050116, AL137558, A08915, AL11022	
AL117626, AF143957, U95114, AL13745	
AC004383, I33392, D44497, AL	
, AL137627, AF207750, A57389, AF11806	
AL133112, U88966, AL117648, AF162270	
AL137488, AC004200, E01614, E13364	
, M96857, A58545, U57352, AL09	
AL133560, AL137537, AL049283, AL117	
AL050393,	-
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AF104032, AF067420, X06	
AL137665, AF169154,	
AL133557, AL137529,	
, AF118094, A65341, U58996, AL08007	
524, A58523, AL133104, AL133067, AL13307	
06743, U78525, AF115392, AL080126, AL13755	
, X72889, AF113677, I48979, U66274	
, U87620, AL133014, AJ005690, AF18221	
, AL137429, AF175903, AF065135	
5, U72621, AL080163, AL122121, A089	
5340, X79812, AL133080, AL133081, AF19259	
AL137539, AL122110, AF100931, AL137	
0, I89934, AL117432, AL133565, AL133	

1584	HSYAP76	876701	Preferably excluded from the	AW411543, AL039599,	, AI351337, AI826980,
			present invention are one or more	AA160380, N67961, A	N67961, AI378493, AI951298, AI090558,
			polynucleotides comprising a	AI348126, AA478324	, AI200956, AA644040,
			nucleotide sequence described by	AW024189, AA587243	, AI812050, AI362845, F29594,
			al	AA776518, AA789114	, AA931516, AI003566,
			is any integer between 1 to 484 of	AI707494, AA970343	, H11327, AA947278, AA076341,
			SEQ ID NO:1584, b is an integer of	AA915984, AI299557	, AW299825, AA024520,
			15 to 498, where both a and b	AA258801, AA169301	, AA342232, AA484880, W90755,
			correspond to the positions of	AA516277, AI015269	, R53617, AA113377, AI379669,
			e residue	AA829839, AA876766	, H05518, AA053830, AI991853,
			NO:1584, and where b is greater	AA810454, AI766365	, R85352, AA502109, AA922383,
			equal to a + 14.	H09142, AI680956, 1	R69168, AA865843, H85022,
				-	AA215481, R06394, AA524191, AA074146,
					R76047, AA528723, F19676, AA588290,
				N56241, N75886, R2	N75886, R22963, AW090423, AA088341,
				R75873,	508387, N98357, N67304,
				AA749208, AA355684	AA355684, AA258709, R87295, AI192394,
				AA477680, AA765589	AA765589, AI886515, AA302356,
				AA670313, H11756, A	
				AW167222, R51947,	AA307613, AA478268, AA641818,
				AI252414, AI312364	
				AW269098, AW268251	
				AW073865, AI670009	, AI473536, AI538259,
				AW409772, AI307604	, AI433157, AI702073,
				AA838230, AI500061	, AW084056, AI633125,
					, AI872910, AL045500,
				AW020397, AW079432	, AL040184, AI648454,
				AI766348, AL036631	, AW162118, AW051088,
				AI698391, AI915291	, AW088691, AI859991,
_				AI582932, AI872423	
				AI866469, AW238688	, W74529, AI281800, AI690748,
				AI569583, AI432030	, AI610770, N75779, AI538564,
				AW161156, AI683173	, AW089275, AA235825,
				AI623941, AI537677	, AI890907, AI612852,
				AL046595, AI918435	, AL047344, AI884318,
				AI569637, AA579618	, AI868931, AA001397,

			AF104032, M92439, Y10655, AL137283, A65341,
			AUGSUG, EUG/43, ', AL117435, AU37
			AF113699, Z82022, I46765, AF
			,050149, AL133568,
			S78214, AL122110, AL049300,
			AL137478, E02349, AL137459, AL117460, AL050155,
_			U88966, AF100931, AL110196, AL049430, AL137529,
			AL117394, AL137705, AF061573, AL137292,
			AL110159, X60786, AF132676, AL133640, AF061836,
			AL110197, X84990, A93350, AF039138, AF039137,
			X83508,
			X65873, AL137479, AR011880, A18788, A21103,
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			AF061795, AF151685, AL133016, S68736, AF090901,
			AR000496, U39656, AL080110, Y09972, AF090896,
			AF008439, AF098162, AF113013, AF054599,
			AF067728, AL117416, AF153205, A07647, I09499,
			, S61953, X87582, Y1
			AL049382, AL117626, I17767, AJ238278, AL122100,
			AJ003118, AL050146, AL122093, AL050092, X98834,
			AL137463, AF113690, AL117644, X83544, AF111851,
			AL049466, AF090886,
			I42402, U00763, E03348, AF118094, ARQ38969,
			AL137538, AL080074, I03321, X59414, AF139986,
			D83032,
			X62580, AL117583, L13297, A12297, AL122121,
			AL122123, E15569, AL080124, AF119337, AF117959,
			AF113689, AF126247, A65340, U67958, AL137560,
			U67328, AL133081, AF151109, AL117649, E08631,
			AL133072, AL110222, AF079765
1585 HCRMV17	876716	Preferably excluded from the	AI492198, AA381672, W44823, AB002357, D26077
		present invention are one or more	

			polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 714 of SEQ ID NO:1585, b is an integer of 15 to 728, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1585, and where b is greater	
1586	HOEKC59	876719	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 1794 of SEQ ID NO:1586, b is an integer of 15 to 1808, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1586, and where b is greater than or equal to a + 14.	AI436209, AW026035, AI401315, AI446530, AA588136, AI591172, AA497132, AA927681, AA497055, AI951115, AI200036, AW238900, AI493315, AI400504, AI089283, AI925204, AW069539, AA857330, AI191461, AI378670, AA410339, AI472923, AA747530, AA766215, AA234951, AA988960, AA037081, AI246277, AI167513, AA704133, AI080251, AI055948, AA614812, AA130081, AI015171, AI493376, AA235125, AA825222, AA449908, AW206209, AA130080, AA029281, W25810, AA613492, Z44379, AI417639, D82431, AI198426, R23635, Z40312, AW390845, D79780, D79680, R24115, AA455230, AW390828, D63116, AA465608, T10625, W51823, AM8198, AA079425, AW390832, D19792, AA258657
1587	HKCSL28	876722	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 363 of SEQ ID NO:1587, b is an integer of 15 to 377, where both a and b correspond to the positions of	1, AA089740, AB003103 9, AI29932, AI245421, AA8723 1, AA927697, AI244692, AI3788 8, AA917836, AA894628, AI2999 5, M87842, M14079, M87859, M8

			ide residues s , and where b				
1,500	UUCEDAK	36676	than or equal to a + 14.	AT052256	AT126717	AW189938	AA745594
9961	ווורו הזמ		y exercion are one	00	AW070663,	Z99376, AI	-
			leotides comprising a	AI784576,	AW327439,	ω,	
			nucleotide sequence described by	AW276639,	AA835672,	A1608763,	N36799, AW247076,
			the general formula of a-b, where a	AA627848,	AI127547,	AA740916,	AW327258,
			is any integer between 1 to 1472 of	AA166916,	AA568685,	AA828239,	Z99375, AA700740,
			SEQ ID NO:1588, b is an integer of	AW327612,	AA812422,	AA099018,	AA761648,
			15 to 1486, where both a and b	AI051506,	AAS73156,	AI025865,	AA503846,
			correspond to the positions of	AA592898,	AA160273,	AA775540,	AA451628,
		_	e residue	AI185757,	AA768416,	AA687268,	AI37.1140,
			NO:1588, and where b is greater	AI371046,	AA074799,	AW029151,	AW250428,
			than or equal to a + 14.	AI138225,	AI089539,	AI004126,	AA809470,
				AI537332,	AI073676,	AI190076,	AI278484,
				AA167073,	AA127406,	AA649193,	AA721424,
				AA715174,	AA978034,	AA524391,	AI923795, W88636,
				AA393865,	AW403551,	AA173982,	AW362155, W73908,
				AI635344,	AA856908,	AA962673,	AI024400,
				AA992622,	AI167830,		AI031946,
				AI752947,	AA100657,	AI922493,	H83589, AA593126,
		•		AA888675,	R54097, A	R54097, AA031733, AI033288,	033288, AA506081,
				AI380802,	AI491801,	AI953284,	AA085335,
				AA127405,	AA515785,		AA076411,
				AA075012,	AA305905,		AI039462, AA450223,
				AA112634,	AA082732,		AW341032, AA725074,
				AA074990,	AA009468,	AA889213,	AA565437,
				AW079297,	AA099096,	AI064753,	AI064753, AA027240, H00352,
				AA173626,	AI380804,	W88554, AA076267,	.076267, AW105351,
				AA076266,	W52167, AW021312,	Æ	AA693887, AA164763,
				AI249663,	AA031732,	AA031732, AA403080,	R89292, R51433,
				AW327440,	H02543, N	52907, AA11	H02543, N52907, AA113337, AA127505,
				AI282747,	AA164762,	AA411811,	AI459951,
				AA133539,	AA514558,	AI197787,	72
				AW393147,	AA314358,	AA933718,	C00036, AA639385,

					H02544, AI696072, H7	AA361575,
						R54151, AA588847,
				W73014, R9	R99520, R89293, AA969406, AI797	39406, AI797468,
				AA864670,	AI083791, AA628031	AA628031, AA974650,
				AA053334,		AI380120, AA058648, T27975,
				AA393799,		AA076505, H94038, AI126113,
				AW449655,	AI686294, T47873,	T47873, T73141, R16766,
				AA810517,	T74664, R07722, R0	R07723, AI300209,
				N45959, H4	N45959, H47972, AI379137, AA903779, AA876048	1903779, AAB76048,
				AA320546,	AA922980, AA782268	AA320546, AA922980, AA782268, R10017, AA644180,
				R15278, AA356761,	1356761, AI688217,	AI688217, R93621, AI476203,
				AI267797,		AA910612, AI201954, R09847,
				AW364121,	AA179728, H47662,	H47662, AW104377, AA872213,
				AI718364,	AI718364, AW166745, AA191273, AA492543,	, AA492543, T83787,
				W24030, AM	AW197934, T11052, AI686637,	.686637, AW351540,
				N55602, AA	AA127491, AA665178, W63552,	W63552, AI143483,
						5216,
	_		_	F06634, T1		36, X73836,
				AL031668,	AC007934, AF076927	
1589	HWBBS84	876726	Preferably excluded from the	AA775676,	AA306997, AW299505	, AA295175,
			present invention are one or more	AI660377,	AI698467, AI925518	
			polynucleotides comprising a			
			nucleotide sequence described by			
			the general formula of a-b, where a	-	-	
			is any integer between 1 to 984 of			
			SEQ ID NO:1589, b is an integer of			
			15 to 998, where both a and b			
			correspond to the positions of			
			nucleotide residues shown in SEQ ID			
	- 1/2		NO:1589, and where b is greater			
			than or equal to a + 14.			
1590	HSIFZ22	876728	Preferably excluded from the	AI554023,	AI913274, AW383970	i, AW383965,
			present invention are one or more	AW383954,	AI539770, AI609013	, AL043107,
			polynucleotides comprising a	AW383974,	AW383967, AW167072	, AW383980,
			nucleotide sequence described by	AIS91170,	AA001432, AI612801	, AW129469,
			the general formula of a-b, where a	AI799420,		, AI97863

			is any integer between 1 to 2108 of SEQ ID NO:1590, b is an integer of 15 to 2122, where both a and b	AW383979, AI375787, AI129128,	AW380739, AA888783, AI073851,	AI289788, AI560125, AI818814,	AL041919, AW383982, AA157885,	
			correspond to the positions of nucleotide residues shown in SEQ ID	AA157573, W67551, D2	AW365658, 9067, AA14	R53920, AV	AA157573, AW365658, R53920, AW363206, AI59001 W67551, D29067, AA143454, AI273137, T29043,	590019, 43,
			NO:1590, and where b is greater than or equal to a + 14.	AI681062, AI648445,	AA862112, C00135, D2	AW383985, 19068, AIS6		609506, 80,
				N74341, AW AW383976,	1189660, AP AW363205,	N74341, AW189660, AA143453, AI168413, AW383976, AW363205, AW392754, T25083,	D2	D29362, L34155,
				X84900, X6	4013, X840)14, U6126	X84900, X84013, X84014, U61261, X85107,	X85108
1591	HCRNB80	876731	Preferably excluded from the	AI750182,		S79910, U37431,	131, S79869	
				AC004079,	264816			
			otides comp					
			nucleotide sequence described by					
			the general formula of a-b, where a					
			is any integer between 1 to 515 of					
			SEQ ID NO:1591, b is an integer of					
			15 to 529, where both a and b					
			correspond to the positions of					
			nucleotide residues shown in SEQ ID					
			NO:1591, and where b is greater					-
			than or equal to a + 14.					
1592	HTPAY47	876732	Preferably excluded from the	AL045837,	AW290917,	AI925409,	AW168903,	
			present invention are one or more	AW068826,	AI083568,	AW026383,	AW262903,	
			polynucleotides comprising a	AI926513,	AI979214,	AI890598,	AI750592,	
		•	nucleotide sequence described by	AW339074,	AA418236,	AW029483,	AW022107,	
			the general formula of a-b, where a	AW295181,	AA664461,	AI752803,	AI740606,	
			is any integer between 1 to 1202 of	AI147688,	AA970819,	AW068765,	AI473816,	
			SEQ ID NO:1592, b is an integer of	AI751522,	AI925816,	AI459360,	AI752768,	
			15 to 1216, where both a and b	AI752291,	AA639417,	AI460028,	AI752525,	
			correspond to the positions of	AI750945,	AI694639,	AA599476,	AW131293,	
٠			nucleotide residues shown in SEO ID	AA242752,	AI750659,	AI889686,	AI888426,	N71781,
			NO:1592, and where b is greater	AI357766,	AW021892,	AI755098,	AA350793,	
			than or equal to a + 14.	AW067910,	AA853461,	AA298896,	AI784082,	
				AA853579,	AA852453,	AA852454,	AA853800,	
				AA307755,	AI925501,	AW021059,	AA976657,	

				AW150473, AW166734, AA627471, R30650, AI752649,
				C01914, AL049389, AL109718, AB033025, I95744, AR053539
1593	H2LBA37	876743	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 675 of SEQ ID NO:1593, b is an integer of 15 to 689, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1593, and where b is greater than or equal to a + 14.	, AA314510, AF121.164
974	HWLFF86	876744	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 932 of SEQ ID NO:1594, b is an integer of 15 to 946, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1594, and where b is greater than or equal to a + 14.	AF121164, AA863031, AA877523, AA741216, AA568880, AW272162, AW135907, AA887896,
1595	HGBAM79	876745	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 861 of SEQ ID NO:1595, b is an integer of 15 to 875, where both a and b correspond to the positions of	AA424088, AA419164, AI003828, T28640, H69474, Y00291, M96023, S56660, X07282, AF110730, AF110729, AF157483, X59473, I09352, I09359, S63196, X57340, X57339, X56674, X57341, M96022, I09358, M96021

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	AI346365, AA641709, AA627539, AI340146, AI909720, AA555216, CI6952, AW014754, AA857163, AA975933, T29526, AI431323, AI269804, AW371982, T61465, D29449, AW268543, M30704, AR052268, M30699, M30703, AR052271, M30698, AR052272, M30700, Y09830, M30701, M30702, AR040760	AA775705, AW361849, AW370643, AW361561, AW378535, AI831033, AA088652, AA968933, AA524822, AA043825, AI620610, AI906062, AI620610, AI906062, AW385411, AW385415, H75542, AW385929, N8, AW379467, AL135407, AM3744235, AI752870, AC006316	AIS71948, AA308400, AA573793, AA314326, AA568312, AA614579, AI925552, AA307578, AA507595, AA614409, AA314825, AA578674, AA582084, AW009769, AA514776, AA588034, AW004668, AA587613, AA858276, AW050700, AI624586, R83818, AI001051, AI910275, AW050690, AA864309, R83377, AA524242, AA507418, AI202532, AI307407, R55389, AI970839, R55292, AI909751,
nucleotide residues shown in SEQ ID NO:1595, and where b is greater than or equal to a + 14.	preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 1243 of SEQ ID NO:1596, b is an integer of 15 to 1257, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1596, and where b is greater than or equal to a + 14.	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 927 of SEQ ID NO:1597, b is an integer of 15 to 941, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1597, and where b is greater than or equal to a + 14.	preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 491 of SEQ ID NO:1598, b is an integer of 15 to 505, where both a and b
	876747	876750	876752
	HKAFU85	HNFEO67	H2MBA27
	1596		1598

			correspond to the positions of	AI910083,	AI909772,	AA614539,	AI909749,
		_	nucleotide residues shown in SEQ ID	AA506787,	X00474, X	AA506787, X00474, X52003, E02904,	M.1.2
			NO:1598, and where b is greater	E03953, X(05322, X05	953, X05322, X05321, X05030	0
			than or equal to a + 14.				
1599	HWLMB30	876753	Preferably excluded from the	AI307407,	AI571948,	AI909772,	AI909751,
			present invention are one or more	A1909749,	AW009769,	AI970839,	AW050690,
			polynucleotides comprising a	AW050700,	AA524242,	AA587613,	AA858276,
			nucleotide sequence described by	AI202532,	AA507595,	AW004668,	AA514776,
			the general formula of a-b, where a	AA578674,	AA573793,	AI925552,	AA614409,
			is any integer between 1 to 266 of	AA614579,	AA588034,	AA308400,	AA582084,
			SEQ ID NO:1599, b is an integer of	AA307578,	AI001051,	AA568312,	R83377, AI624586,
•			15 to 280, where both a and b	AA314326,	AA314825,	AA507418,	X00474, X05322,
			correspond to the positions of	M12075, X52003,	E02	904	
			nucleotide residues shown in SEQ ID				
			NO:1599, and where b is greater				
			than or equal to a + 14.				
1600	HHEBN60	876760	Preferably excluded from the	AI131324,	AL037422,	AL037391,	AW161774,
			present invention are one or more	AI890947,	AA122289,	AA584305,	AW273236,
			polynucleotides comprising a	AI862040,	AW085692,	AI209167,	AA148506,
			nucleotide sequence described by	AI351762,	N66647, A.	I523188, AM	N66647, AI523188, AW273178, AI830451,
			the general formula of a-b, where a	AA452008,	AA705906,	AL043832,	AI571577,
			teger between 1 to 1519	AI219060,	AI361659,	AA632645,	AA662786,
			SEQ ID NO:1600, b is an integer of	AW273354,		AI885486, AA627153,	AI050005,
			15 to 1529, where both a and b		W56473, A	W56473, AI266655, C75555,	75555, AA884431,
			correspond to the positions of	W70047, W	70048, N634	491, N64411	W70048, N63491, N64411, AW055257,
			de residues s	AI424319,	AI554547,	AI554547, AI521110, AI559699	AI559699,
			NO:1600, and where b is greater	AI623228,	N92821; AA160261,	4160261, AF	AA135865, AA171948,
			than or equal to a + 14.	AI619980,		AA169427, AI434909,	AI434909,
				AW021267,	AI539602,	N94794, H03661,	3661, AA999936,
				C17025, A.	1055978, H(C17025, AI055978, H03756, AI567074,	57074, AA151579,
				AI918516,		H88943, R70308,	70308, AI904987,
				AA345034,	AI970814,		70632, AA135864,
		_		AA740380,	AA156595,	AA353886, R22230	R22230, AA618325,
		_		D56914, H4	44681, AI35	H44681, AI355451; AI955112,	55112, AI919589,
				C75412, A			AI907423, T50659,
				AW263380, D56915,		C02126, AI28445	34452, R31847,

				T40470, AI904794, AA384278, AI568036, T39196,
				C75672, T27972, D55752, R22288, AA862190,
				AI907464, AA149395, AA513034, R35775, AA484012,
				AA649723, AA160260, AA074934, AA262411,
				AA828667, AA501402, AW302880, AI076612,
				, AA975564,
1091	89ОМЭОН	876762	Preferably excluded from the	AA603949, AI680975,
			present invention are one or more	AI393833, AI770102,
			polynucleotides comprising a	
			nucleotide sequence described by	AI885125, AI373081, AI580943, AI393771,
				AA749301, AW338708, AI250780, AA287845,
			is any integer between 1 to 3082 of	AW453050, H71837, W03966, AA152044, AA603836,
			SEQ ID NO:1601, b is an integer of	AA287846, AA042955, N99630, W02451, N25637,
		•	15 to 3096, where both a and b	AI917997, AA244066, R63787, AA578977, AW239000,
			correspond to the positions of	R78310, H54574, AA037115, N34235, AI240141,
			nucleotide residues shown in SEQ ID	AW130305, H02870, AA042815, R73884, AA334992,
			NO:1601, and where b is greater	
			W	R82819, AI128764, R63733, AA664138, AA953035,
				AA113801, R63857, AA298118, R23143, R62758,
				T69806, AA303428, R34175, R73971, H59544,
				R23144, T70792, R31823, R82820, AI933547,
				AA742952, AI453225,
				AA327996, AW338192, R22283, R77939, AI240290,
				N72673, N95485, AA152084, AI383282, H60415,
				N98505, AW361055, R32084, R31777, R34297,
				R32031, AA374818, AA300327, AI076967, AA622059,
	-			R63858, N73903, AW150955, AI368478, AA037154,
				AW087179, AL080209, X67780, AF130561, M96248,
			:	M64474
1602	HHFCP36	876764	Preferably excluded from the	AA347863
			present invention are one or more	
			polynucleotides comprising a	
			nucleotide sequence described by	
			the general formula of a-b, where a	
			is any integer between 1 to 322 of	

			SEQ ID NO:1602, b is an integer of	
			15 to 336, where both a and b	
			nucleotide residues shown in SEQ ID	
			NO:1602, and where b is greater	
			than or equal to a + 14.	
1603	HTXKH86	876767	Preferably excluded from the	AA314774, AI291017, AA191539, AI298290,
			present invention are one or more	AA147791, AW238920, AA308544, AA187762,
			polynucleotides comprising a	AA081307, AA075926, AA773549, W52392, AA780574,
			nucleotide sequence described by	AL038991, AA307244, AA181578, AA081167, C06415,
			the general formula of a-b, where a	AW402249, AA165319, AA132481, AW247110,
			is any integer between 1 to 1021 of	AA076454, AA079384, AA304499, AA181561,
		_	SEQ ID NO:1603, b is an integer of	AI857405, T35498, C06389, AA181655, AA314234,
			15 to 1035, where both a and b	AA352654, Z45227, AA992505, AW000888, AI651014,
			correspond to the positions of	AI392985, T34265, AI344273, AW341319, AA190808,
			nucleotide residues shown in SEQ ID	R71708, AF104669, U87954, AR035973, U59435,
			NO:1603, and where b is greater	X84789, U43918, U50137
			than or equal to a + 14.	
1604	HISCI72	876771	Preferably excluded from the	AI743600, AI885169, AI937505, AI042181,
			present invention are one or more	AA854952, AIS22015, AA400219, AI522002,
			polynucleotides comprising a	AA305093, N26064, AI888285, AA400130, AW296334,
			nucleotide sequence described by	AW292016, AW440393, AI146794, AA187458,
			the general formula of a-b, where a	AI262079, AA855005, AI476446, AA187590,
		_	SEQ ID NO:1604, b is an integer of	_
			15 to 2231, where both a and b	AW163208, AW070692, C06284, AA838476, Z43206,
			correspond to the positions of	C05759, AA190468, AI680041, AA635314, AI034110,
			nucleotide residues shown in SEQ ID	AA622708, AI000051, R64675, W44694, D60048,
			NO:1604, and where b is greater	AA805958, F07813, Z40908, AA565995, F02659,
			than or equal to a + 14.	AI471921, F05522, F05523, AI034108, R27644,
				AW236720, AA039917, AW163735, R64676, R27550,
				W38645, F01794, F01795, AW263460, D52614,
				AW151942, AA090824, C00912, X92396, AJ225782,
				X96737, AJ004799, AJ225808, X95807, AJ133541,
				AJ133539, AJ225807, X95806
1605	HJACJ75	876773	Preferably excluded from the	AA309052, AW247981, AA311506, T87086, AA352616,

			present invention are one or more	AW339919, R01803, AW054854, H63371, AI09755	1097555,
			polynucleotides comprising a		2,
			nucleotide sequence described by	AW392909, H45736, U18300	
			. formula of a-b,		
			is any integer between 1 to 665 of		
			SEQ ID NO:1605, b is an integer of		
			15 to 679, where both a and b		
			correspond to the positions of		
			.0		
			NO:1605, and where b is greater		
			Ψ		
9091	HTEDS58	876776	Preferably excluded from the	AA147098, AA506483, AA459122, AA553631	7,
			present invention are one or more	AA687219, AA639000, AA507321, AI475344	4
			polynucleotides comprising a	N47467, H15303,	W69943,
			nucleotide sequence described by	AA419435, W69833, AA680161, T27895, AI	AI680311,
			the general formula of a-b, where a	H93979, C75158, H93980, R25544, AA223335,	335,
			is any integer between 1 to 1663 of	AI758259, AW079484, F02620,	AI933243,
			SEO ID NO:1606, b is an integer of	2, F02623, AI191766, R12384,	AA371184,
			15 to 1677, where both a and b	AA714796, AI383543, T69739, R09794, AI	AI873805,
			correspond to the positions of	AI581822, AI371311, R15273, AA093267, AA312224	AA312224,
			nucleotide residues shown in SEQ ID	S67325, X73424, AB000886, M14634, M13573,	573,
			NO:1606, and where b is greater	AJ006497, AJ006496, AJ006499, AJ006494,	4,
			than or equal to a + 14.	AJ006488, AJ006491, AJ006493, AJ006492, M31167,	2, M31167,
				٠.	31168,
				AJ006489, AJ006490	
1607	HUVHP60	876789	Preferably excluded from the	AA347492, AA307478, R18976, AA233030	
			present invention are one or more		
			polynucleotides comprising a		
			nucleotide sequence described by		
			the general formula of a-b, where a		
			is any integer between 1 to 1195 of		
			SEQ ID NO:1607, b is an integer of		
			15 to 1209, where both a and b		
			correspond to the positions of		
			nucleotide residues shown in SEQ ID		
			NO:1607, and where b is greater		

			than or equal to a + 14.					
8091	HUFC129	876791	Preferably excluded from the	AW007623,	AI963511,	AIS87104,	AI453405,	
			present invention are one or more	AI694729,	AI796832,	AW363443,	AW387811,	
			polynucleotides comprising a	AW387793,	AI826957,	AW361899,	AI955696,	
			nucleotide sequence described by	AI955780,	AI827005,	AW387799,	AI828295,	
_			the general formula of a-b, where a	AW192552,	AA581220,	AA527188,	AW387817,	
			teger bet	AW363244,	AI818260,	AI956167,	AI801443,	-
			SEQ ID NO:1608, b is an integer of	AI904486,	AI400372,	AI921063,	AW338519,	
			15 to 2608, where both a and b	AI693877,	AI074261,	AI927711,	AI956102,	
			correspond to the positions of	AI920992,	AI972695,	AI911695,	AI828218,	_
			de residue	AW076111,	AI682785,	AI921387,	AW387812,	
			NO:1608, and where b is greater	AW337936,	AW363218,	AW364488,	AI346975,	-
-			than or equal to a + 14.	AI913862,	AW440967,	AW130304,	AW360772,	
				AI696946,	AI672948,	C05920, AI	.587485, AW070932	0932,
	_			AI635943,	AI262029,	AI739440,	AA100719,	
				AI955836,	AI262264,	AW376483,	AW130542,	
				A1972967,	AW175800,	AW387796,	AA579753,	
				A1446049,	AI569938,	AI934313,	AI609930,	•
_				AI677998,	AI431963,	AA553880,	AI828330,	
				AI597812,	AA040073,	AW360835,	AA917638,	
_				AW377104,	AI682718,	AI354639,	AW376508,	
				AW192548,	AI962102,	AW376484,	AW392307, U4	U47705,
				AI813978,	AW362727,	AW361642,	AA828073,	
				AI261531,	AI277071,	AW136050,	AW361304,	
				AI934325,	AA152037,	AI695028,	AI631388,	•
_				AW377034,	AA316326,	AI470301,	AI962061,	
				AW377083,	AW360762,	AW362547,		
				AW391349,	AW375920,	AW376475,	AW243579,	
_				AA130547,	AW365061,	AI961867,	AA135037,	
				AA581264,	AI250167,	AI453469,	AI696953,	
				AW376234,	T29561, A	AI589481, AI	AI582988, AW38771	7713,
				AI537547,	AW387715,	AW376010,	AI926514,	
				AA132781,	D45505, AA	D45505, AA367446, AA	838269,	AA295348,
				AI828399,	AI473526,	AI587351,	5, T	93569,
				AW376489,	AW393447,	AI584131,	AA132182,	
				AW360942,	AL121028,	AI569894,	AI264699,	

ALS64753, AW371781, AA587700, AW387798, AB58790, AB587700, AW373707, AA584940, AI872586, AW373781, AW373781, AA053542, AW374712, AA053542, AW374712, AA13613, AW373627, AA366104, T29474, AI US4607, AW373640, AW37372, AW37372, AW37372, AW37364, AW33366, AW37372, AW37372, AW37364, AW337364, AW37372, AW37366, AW373766, AW37372, AW37366, AW373766, AW37372, AW37376, AI686197, AW29776, AI686197, AW297144, AI763361, AW29776, AI686197, AM29776, AI686197, AW29776, AI6861936, AW29776, AI6861936, AW29776, AI6861936, AW29776, AI681936, AM29776, AI681936, AM29776, AI681936, AM29776, AI681936, AM29776, AI681936, AM29776, AI6861936, AM29776, AI686197, AM29776, AI6861936, AM29776, AI686, AM29776, AI686197, AM29776, AI686, AM29776, AI686, AM29776, AI686, AM29776, AI686, AI686, AM2976, AI686, AM2976, AI686, AI686, AM2976, AI686, AM2976, AI686, AI686, AM2976, AI686, AI686, AM2976, AM29776,	AL264/53, AW387/102, AM387806, AM05809, AAS84940, AI872586, AW176585, AW364936, AW373781, AW373707, AA834430, AW374782, AA584940, AI872586, AW176585, AW364936, AW373781, AW373783, AW37836, AIG11749, AA013542, AW374412, AW380829, AW375785, AA366104, T29474, AI991653, AW364960, AW365104, T29474, AI991653, AW364960, AW373705, AW364960, AW373706, AW36502, AW376976, AW36502, AW37691, AW373706, AW36502, AW37691, AW37091, AW37091
from the re one or more orising a described by of a-b, where a sen 1 to 1999 of s an integer of	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 1999 of SEQ ID NO:1609, b is an integer of
from the re one or more orising a described by of a-b, where a sen 1 to 1999 of s an integer of	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 1999 of SEQ ID NO:1609, b is an integer of
from the re one or more orising a described by of a-b, where a sen 1 to 1999 of s an integer of	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 1999 of SEQ ID NO:1609, b is an integer of
Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where is any integer between 1 to 1999 SEQ ID NO:1609, b is an integer of the general sequence of the	
	876795
HCRNO02	

		15	15 to 2013, where both a and b	AI367070,	AA976607,	AA583461,	AI249930,
		ပ္ပ -—	correspond to the positions of	AW051844,		AA507715,	AI954585,
		nu	nucleotide residues shown in SEQ ID	AA922244,		AA126244,	AI087863,
		NO	NO:1609, and where b is greater	AI251918,	AI334712,	W67736, AI242730,	242730, AA101742,
		th	than or equal to a + 14.	AW135527,			AI347209,
				AI286337,	AIS81372,	AI469691,	AA069014,
				AA934842,	AA508884,	AI887809,	AA831979,
-	-			AI244186,	N50480, AI	275702, NS	N50480, AI275702, N50424, AA736752,
				C20724, NS	5586, AW30	4156, AA45	C20724, N95586, AW304156, AA459318, AW192272,
				AI275964,	AA947333,	AA902224,	AI220977,
				AA742300,	AA321817,	AA553858,	R63954, AI933896,
				AI569580,	AW084360,	AI802071,	AA888637,
				AI802496,	AA364540,	AA330481,	AI623357,
	_			AA459100,	AI879891,	AA321816,	AA806651,
				AW270487,	AW117230,	N73503, AI	AW117230, N73503, AI763427, AI570080,
				AA602961,	T27344, W2	5008, AA30	T27344, W25008, AA306002, AW377570,
				AA016984,	W67735, AA	377036, AA	AA377036, AA092406, AA876851,
				R27168, AA069079,		U18914, S820	S82081, U35428, D8257
1610 HAU	HAUAF56 876798	_	Preferably excluded from the	AA843663,	_	AI652163,	AI741572,
		pr	present invention are one or more	AI734839,	AI191667,	AI311840,	AI092011,
		Od D	polynucleotides comprising a	AA838667,	AI651387,	AW236921,	AW241575,
		nu	nucleotide sequence described by	AA861653,	_	AA602368,	AI689816,
	_	t t	the general formula of a-b, where a	AW051840,	AI354951,	AA573089,	AI148406,
		is	is any integer between 1 to 590 of	AI141828,	AI183782,	AI194006,	AI693445,
_		SE	1:1610, b is an	AI635512,	AI493869,	N90872, AW	N90872, AW237388, AA126737,
		15	15 to 604, where both a and b	AA732844,	AI192168,	AI217045,	AA137055,
		8	correspond to the positions of	AA994789,	AI493086,	AA845631,	AI094429,
		nu	nucleotide residues shown in SEQ ID	AL047557,	AA181124,	AI140430,	AI860338,
	-	8	NO:1610, and where b is greater	AA723326,	AA506514,	AI718897,	AI142056,
		th th	than or equal to a + 14.	AA694462,	AA527690,	AA719919,	W60495, AI128784,
		_		AA295736,	AA719929,	W74729, AA	AA046090, AL079932,
						AA777211, AA	AA187497, W60781,
				W02217, AL	AL047558, AI	AI962738, WS	W57590, W58378,
				AI040455, N78658,			AI092598, AI127083,
				AI767352,	C00790, AI	AI796294, F2	F21069, AI962745,
				W58054, R8	R82964, AI127007	7007, AA319961	9961, H25260,

				AA046133, F29476, AI024494, D57900, AA187496,
				R27633, F15904, N92901, F16228, AI880466,
				AA513941, AI028160, AA320194, AI942291, W15147,
				AA515161, AA319909, H27992, AA137126, AA032269,
				W17092, AA305767, AA317925, AA315585, AA316680,
				AA385920, AA082685, AA393514, AA319917, R82782,
				W21107, H58270, W60536, AW385090, AI857611,
				H74142, W23645, F37285, AI831575, AW009545,
	_			AI907307, F00610, N86957, AI955298, AI904744,
				31730, AA30
				AA314317, AW131256, AW173066, AI590946,
		_		AI880624, AI566275, N91884, AI610714, AA640156,
				AA804541, AI638798, J02874, A98023, M94856,
				AF181449, AF102872, AF136241, AP000547,
				AP000365, I88901, R82963
1611	HHEUM25	876802	Preferably excluded from the	AI817822, AA148948, N50594, N25959, AA086480,
			present invention are one or more	
		_	polynucleotides comprising a	AA160920, N50540, AA602221, AA160014, H53938,
		_	nucleotide sequence described by	AI079093, AI015698, AI439431, T89890, AA086479,
			the general formula of a-b, where a	H83411, AB033097
		_	is any integer between 1 to 965 of	
			SEQ ID NO:1611, b is an integer of	
			, where b	
			correspond to the positions of	
		_	nucleotide residues shown in SEQ ID	
			NO:1611, and where b is greater	
			than or equal to a + 14.	
1612	HWLQW0	876804	Preferably excluded from the	, N67220, AI538999, AW119213,
	∞		present invention are one or more	N91158, AI
			polynucleotides comprising a	AI369016, AI091413, AI435427, AW296026,
			nucleotide sequence described by	AW195056, AI765593, T16459, H99837, R55315,
			the general formula of a-b, where a	D29082, H88285, AI537645, R33635, D63011,
			is any integer between 1 to 490 of	AIS53628, AI923565, AI270171, H49679, D61792,

			SEQ ID NO:1612, b is an integer of	H52824, R55417
-			15 to 504, where both a and b	
			correspond to the positions of	
		-	nucleotide residues shown in SEQ ID	
			than or equal to a + 14.	
1613	HOEOP07	876807	Preferably excluded from the	AI290876, AI765569, AI808777, AI338031,
			present invention are one or more	AA913566, AA573434, AI568487, AW175945,
			polynucleotides comprising a	, AW41876
			nucleotide sequence described by	, AA68795
			the general formula of a-b, where a	
			is any integer between 1 to 1636 of	AA315078, AI802767, AA581469, AA620711, H45711.
			SEQ ID NO:1613, b is an integer of	
			15 to 1650, where both a and b	AI809670, AW008766, AI915360,
			correspond to the positions of	AI687397, AW023240, H45668, H04001, AA297249,
			ന	0, AW188056, D25944, AW196645, AAS06
			NO:1613, and where b is greater	70465, AI784132, AA382289.
		_	than or equal to a + 14.	AI537449, D58213, AA298492, AA29880
				A904960, AA298494, AW020800
				, AF105036
1614	HCQAE79	876809	$\frac{1}{2}$	AI346844, AW001371, AI991265, AI246778,
			present invention are one or more	2, AI832475,
		_	ö	AW00080
	_		nucleotide sequence described by	, AI983400, AI673613,
			al formula of a-b, when	AI991308, AA857748, AI672894, AI732375,
			teger between 1 to 973	AI73235
			:1614, b is a	AW001307, AA32745
			15 to 987, where both a and b	AI688199,
			correspond to the positions of	\sim
			nucleotide residues shown in SEQ ID	
	-		NO:1614, and where b is greater	
			than or equal to a + 14.	
1615	HCQDR53	876811	Preferably excluded from the	AI738919, AI923216, AW237190, AI769620,
			present invention are one or more	AI905420, AI905431,
			polynucleotides comprising a	

			nucleotide sequence described by	AA774861, T85091, AA150805, AA666115, AA150811, T33125, AA173650, T84156, R49735, AA150702,
			teger between 1 to 1473 o	T35291,
			SEQ ID NO:1615, b is an integer of	Z39956, AA150709, F03307, R48157,
			15 to 1487, where both a and b	T35290, R40351, T35286, H71220, F03153, D61519,
			d to the positions of	AI650460, H71219, AF034745, AF034746
			nucleotide residues shown in SEQ ID	
			v	
9191	HOEFO36	876816	딚	AI453687, AI571506, AI417180, AI453138,
			present invention are one or more	AA993886, AL048366, AI587024, AA769711,
			polynucleotides comprising a	AA906543, AI333633, AI692876, AW007640,
			nucleotide sequence described by	AI399951, AI983818, AI750469, AI433964,
			the general formula of a-b, where a	AW130422, AI355200, AI567515, AW069544,
		_	is any integer between 1 to 699 of	AI367996, AW338539, AI925385, AI583403,
		-	SEQ ID NO:1616, b is an integer of	AI014460, AI077522, AI435310, AI969659,
			15 to 713, where both a and b	AA149832, AI016334, AI016317, AI804042,
			correspond to the positions of	AW068411, AA131691, AI339632, AI750268,
			nucleotide residues shown in SEQ ID	
			NO:1616, and where b is greater	AI247016, AI338848, AW073799, AI753153,
			than or equal to a + 14.	AW068385, AI378389, AW073223, AI752287,
				AA600284, AI474336, AI359229, AA569973,
				AI342311, AI623621, AI753719, N23207, AI587013,
				AA149811, AA723444,
				AI016443, AI961932,
				AA252895,
				AI635286, H88017, AW296238, H38240, AA131706,
				88729, AI25
				AA194241, AI520853, AW068232, AI566383,
				AA853382, AA055161, AI610126, AW021156,
				AW021155, AI359367, AA586748, H78023, T79480,
				AA055064, T94348, AI033179, AA677178, AA976366,
				, AA156786, AA131536
				T28255, AI701212, R40533, C16582, C21348,

				D25653, H88728, L12350
1617	HFIAL22	876817	Preferably excluded from the	AI346330, AA149866, AW190828, AA149859,
			present invention are one or more	ω,
			polynucleotides comprising a	10126,
			nucleotide sequence described by	AI247016, AI753179, AI160032, AA476585,
			the general formula of a-b, where a	9, AI130835,
			teger between 1 to 3508	AI016334, AI378389, AA600284, AW376487,
			SEQ ID NO:1617, b is an integer of	3, AI804042,
			15 to 3522, where both a and b	AW068385, AA677178, AA435731, AI750719,
			correspond to the positions of	AI752286, AW376482, N23207, AI075364, AI623621,
	-		de residues s	AI359367, AI752287, AW068222, AI587013,
•			NO:1617, and where b is greater	
			than or equal to a + 14.	AI246892, AI753719, AW073223, AA252872,
				AI417168, AI955590, W19516, AA397612, AA137054,
				AA316564, W94600, AI750531, AA723444, AI453687,
				AI571506, R51145, H88729, AA331775, AA313295,
				AA481319, H78022, AA307252, AI351084, AA316570,
				AA625464, AW023185, N83257, AA448908, R14334,
				0, AI520853, AI566383,
		_		AA639814,
				, AA853653, W86005, AW
				, AA055064, AA906543
		_		6, T94703, AI333633, AI
				9, AI453138,
		_		, AI692876,
				0
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				1,
				4, AI750269,
				AI583403, AI567515, W46226, AI969659, W46227,
•				
				AA149811, AA131691,
				W068131, AI635286,
				AI783830, AI961932, N66997, AI016443, H88017,

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AI913843, AI445548, L12350, M81339, X96540, L07803, M60853, M87276, M64866, X87620, M62462	7, AW051723, AA933088, AI24604	AI702461, AA612941, AAU17379, A1362464, AA173916. A1474790. AI802234. AI863510.	, AI284788, AA724009,								AI140351, AI859347, AA530873, AA121548,	AI815642, AA768342, AI864674, AA127712,	AA722381, AA987515, AW275917, AA417302,	AI354682, AI025466, AI859814, AA130959, N92869,	AA100477, AW190165, AA768339, AI920875,	AI051671, AW089493, AA417265, AA587755,	AA045598, N21328, AA314322, AI371694, AA844332,	AA043186, AI567303, R83064, AI350331, AW193146,	AA580315, AI039892, AA828283, AI952434,	AW377665, AI289086, AA100476, AI014387,	AA917482, AA975893, N21020, AA621534, AA045597,	H94056, AA306867, AW406948, AIS64973, AI816957,	AA729835, AI289415, AW103201, AI187288,	AA661773, H80956, W04309, AW088039, AI018462,	AA649285, AI083853, AI952495, AI419448, N47889,	R89903, N27984, T40562, D82429, N80197,	AA868207, AI955989, AI091426, AI873582,	AW138496, H81296, AI288157, AI833059, T91268,	R63140, AA130829, D12288, AA298770, AI699667,	AI942324, AA310276, W22908, AA074395, D12293,	T91580, AA342276, H81350, AA053266, AA353671,	AI202414, AI832968, AA342277, AW084334, W25596,
	14.	present invention are one or more	nucleotide sequence described by	the general formula of a-b, where a	is any integer between 1 to 888 of	SEQ ID NO:1618, b is an integer of	15 to 902, where both a and b	correspond to the positions of	nucleotide residues shown in SEQ ID	than or equal to a + 14.	Preferably excluded from the	present invention are one or more	polynucleotides comprising a	nucleotide sequence described by	the general formula of a-b, where a	is any integer between 1 to 1144 of	SEQ ID NO:1619, b is an integer of	15 to 1158, where both a and b	correspond to the positions of	nucleotide residues shown in SEQ ID	NO:1619, and where b is greater	than or equal to a + 14.										
	876822										876823																					
	HWLMN8	~									HCGTC91																					
	1618										1619																					

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polynucleotides comprising
nucleotide sequence
the general formula
SEQ ID NO:1620, b is an integer
15 to 2260, where both a and b
correspond to the positions of
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NO:1620, and where b
than or equal to a +
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701, AC004000, AF038458,
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AP000359, AC005874, AF134471, AC005091,
AC002326, AC004216, U93305, AC004985, AC007845
, AC002351
Z98949, AC004865, AL080243, AC005086, AL049795
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AL117337, AL049843,
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AL021546, AF165926, AC005004
AC004531, U52112, AL078602, AC00605
, AC003010, Z93020, AL022320,
AL049869, AP000117,
AL049748, Z97054, AC006390, AC006197, AP000104,
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, AL023553, Z95114, AC006449, AC005
.C002375, AC006160, AC
, AP000513,
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AC004921, Z94721, AC010205, AF073485, AC004257
AL021707, AC005736, AC002364, AC004687, Z97630
AL080317, AC002465, AL035405, AC004858,
AC003037, Z98036, AC000003, AC003108, AC005180
, AC004021, AC004526
0, AC005280, U80017, AC0025
P000014, AB023049, AC004882, AC005839,

876830 Preferably excluded from the present invention are one or more ANIT4828 polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a AIO33267 is any integer between 1 to 1063 of 150000	Is any inceper between 1 to 1063 or h52695, 129050, A1651192, W26286, H92737, SEQ ID NO:1621, b is an integer of H68163, M76180, M88700, M74029, M84601, M84592 15 to 1077, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1621, and where b is greater than or equal to a + 14.	Preferably excluded from the present invention are one or more polymucleotides comprising a nucleotide sequence described by a horse a polymucleotide sequence described by a horse a polymucleotide sequence described by a horse a processor is any integer between 1 to 2363 of AB28256, AB31652, AI139518, AI336313, AA700790, is any integer between 1 to 2363 of AB28256, AB31652, AI139518, AI336313, AA700790, is any integer between 1 to 2363 of AB28256, AB31652, AI139518, AI336318, AI34641, AI141849, AI33644, AI181818, AI33641, AI43641, AI43641, AI14364, AI33644, AI15641, AI14364, AI14364, AI33644, AI1649, AI3364, AI349421, HI5591, AI33266, H50726, H61529, M51822, M51823, M31248, M63122, AI336318, M31248, M63122, AI33631, M31248, M63122, AI33631, M31248, M63122, AI364312, AI36434, AI16591, AI36326, H50726, H15534, AI16494, H15591, AI33266, H80726, H15534, M31268, M31222, AI364312, M31248, M63122, M313248, M63122, M31248, M63122, M313248, M63122, M31322, M313248, M63132, M313248, M63132, M31324, M31322, M313248, M63132, M31324, M31322, M313248, M63132, M31322, M313248, M31323
HCQDG08 8		HE8BX38 8

	AB028451, AF079763, A91160, AL117457, AL137480,
	A91162, AL049423, AL049347, X99226, AL023657,
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	Z97214, A08456, A31057, I33392, A08912, A08911,
	AF060555
	76508, A
	A
	Y10655, A
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	AF111112, A07588,
	1, Z82022,
	AF0615
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	AL137557,
	, AL137574, AL122100,
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	AF02681
	Σ0
	AL137711, AF044323, AL0801
	, I18358, I34395, AF032666,
	9, I89934, AL031346, X61970,
	AL110196, AL04943
	AF116573
	X72889, I77092, AL137537, E12747, A92311,
	A5
	4728, AB026128, AL137476
	I79595, AF002985, AF100781, AL050172, AL110197,

				AF106697, U68387, X01775, AF139373, AL137665,
				X06146, X96540, S61953, A86558, A41575, X00474,
				AL133080, AF076633, AF159615, AF080622, U37359,
				L04859,
				AB019565, A12558, AF113019, AF100931, Y16645,
				U70981, Y11254, AL122050
1623	HMVCR68	876836	Preferably excluded from the	
			present invention are one or more	AA588565, AA424137, AI299200, AI143920,
			polynucleotides comprising a	AA021117, AI913301, AW151208, AA425305, N47966,
			nucleotide sequence described by	AI436446, AI685061, AF052498, AW081049,
			the general formula of a-b, where a	AW084051, AA451690, AW182326, AI332899,
			is any integer between 1 to 1244 of	AA169542, AA169443, AA954593, AA042910,
			SEQ ID NO:1623, b is an integer of	AA455865, AA149424, AI432492, AA460942, N47904,
				AA319689, AI377265, AA042923, AA461248, H20482,
			correspond to the positions of	AI702363, AI371418, H85541, AW351484, AA151489,
			nucleotide residues shown in SEQ ID	AI955508, AA385706, D79614, AA369939, AA834737,
			NO:1623, and where b is greater	AW175964, H50494, AI291715, AI418716, AA861788,
	_		than or equal to a + 14.	AW339974, AA369940, H87923, AA452637, AB033080,
				D42138, AF011794
1624	HFCAI79	876837	Preferably excluded from the	AL048933, AI271440, AI092964, AI741387,
			present invention are one or more	AI760926, AI333315, AI680148, AA889492,
			polynucleotides comprising a	AW190196, AW365955, AL048932, AI416991,
			nucleotide sequence described by	A1923885, A1445890, A1138940, A1687147,
			formula of a-b, where	, AI082757, AA280201,
			is any integer between 1 to 2455 of	, AW079043, AW001900,
			SEQ ID NO:1624, b is an integer of	AW365942, AI079486, AW451587, AI566301,
			15 to 2469, where both a and b	AI623964, AI032887, AW365973, H22632, AI498456,
			correspond to the positions of	AI270190, AW023890, AW137893, N40556, H47810,
			nucleotide residues shown in SEQ ID	AI336798, H52365, AI933592, AA371581, H52364,
			NO:1624, and where b is greater	AA904952, H22633, AA338820, AI537552, R16961,
			than or equal to a + 14.	T82008, H96979, AI565231, AA377237, T81883,
				T71558, R16906, C01340, AI761493, AA280380,
				N46600, H48145, AW021702, AA887860, AA377236,
				T71263, H42623, T71208, AC004849
1625	HBIOH43	876842	Preferably excluded from the	AL049077, Z43264, AA362903, H44830, AA347303,

			present invention are one or more	M23148 A	7 8C198 Z	91666		Γ
			leotides comprising a					
			nucleotide sequence described by					
		·	the general formula of a-b, where a					
			is any integer between 1 to 1267 of					
			SEQ ID NO:1625, b is an integer of					
			15 to 1281, where both a and b					
			correspond to the positions of					
			nucleotide residues shown in SEQ ID					_
			NO:1625, and where b is greater					
			than or equal to a + 14.					
1626	HOEMJ36	876856	മ	AA910951,	AA843679,	AI348072, P	AI125272,	Γ
			present invention are one or more	AI042167,	AA845606,	AW129714, A	AI927609,	
			polynucleotides comprising a	AA868244,	AI978910,	AIS25551, W	W06825, AA843914,	
			nucleotide sequence described by	AA779705,	AW130928,	W61040, W91	W91932, AI831445,	_
			the general formula of a-b, where a	AW247636,	AA186566,	AI359205, A	AA523378,	
				AI186133,	AI160604,	AI041480, A	AI198816,	
			SEQ ID NO:1626, b is an integer of	AI378985,	AI207388,	AA720662, A	AA181832,	
			15 to 1355, where both a and b	AA928300,	AA890438,	AI688759,	AA393736,	
			correspond to the positions of	AA151916,	W73728, A	AI184656, AI4	473972, AW272617,	
			$\boldsymbol{\sigma}$	AA719242,	AA890475,	AA933747, A	AA534300,	
		_	•	AA987916,	AA622766,	AI371055,	AA878593,	
			than or equal to a + 14.	AI811357,	AI829846,	AI246201,	AA987453, N21142,	
				AA191541,	AI345998,	AI142485,	AA307417,	_
				AA393794,	w	AA934733,	AW082787,	
				AW362863,	W96444, A	AI343759, AW073775,	373775, N26594,	
				AI624204,	AI075412,	W73785, AA7	AA706402, AI075444,	
				AA312077,	AW370975,	AI304681, AA305477,	4A305477,	
				AW370958,	AI339961,	AA988926, AI798191,	AI798191, H96572,	
				AI631255,	AA916632,	N21361, AA393864,	393864, AI242708,	
				AI186143,	AI344381,	AI002050, AA829718	A8829718,	<u> </u>
				AA666025,	AI301839,	N31157, T51	N31157, T51961, W96541,	
				AI186650,	AA450264,	N70868, AA1	AA189020, W35262,	
				AI335966,	AA868435,		AI718683,	
				AI285022,	AW380029,		W79062, W56704,	
				AA450265,	AI203443,	AA313952, H	H05891, AA029676,	7

	AI253584, AI750319, W744
	AAS41387, AI915283, AA953221, AI095790,
	AA687834, N63798, H72663, AA627355, N33299,
	W56739, N44829, H10500, AA223727, AW002227,
	AA961262, AW440854, N92556, C17191, AA223815,
	AI750318, H79841, H50961, AA703995, AA305808,
	AA024948, R91859, R96677, W56383, AA332390,
	AW440710, T28956, AA912076, N57269, N92539,
	W94895, R91038, H13004, AA082120, R92698,
	AI932893, AA459672, AA459794, AA189019, T90302
	N78866, H78774, AA361890, N49784, AA305857,
	AA361459, AA765973, AA361675, AA352730,
	AA771826, H72664, N94156, AI613134, N50485,
	AA628033, F02479, C17291, AA063528, R56364,
	AA459660, W39039, AA642158, H62620, AA352976,
	AA628038, AA729743, AA147291, T82974, AI749422
_	AA459783, R57554, AA729543, T52041, AA143387,
-	N39666, W24824, AA742384, W17271, H62547,
-	AA191268, N50430, AI468860, N54292, AW382069,
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	AF106657, AR068753, X72889, AF017437, AF118094,
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	L04504, AF067420, AF104032,
	ALO80110, Z82022, AB025103, X59812,
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	3, X66871, AL049283, AL050190
	3,
	8, AF039138, AF039137, AL0967
	ω,
	U73682, D16301, AI
), AL050208, X81464, AF113694, X875
	AF115392, AF192557, AF153205, AL1
	A08908, AL050024, X63162, AL
	F017152, X65873, AL133560, AF1111
	I26207, AL117416, AF151109, Z
	U72621, AL
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	, AJ006417, AF102578, x
	89934, AB030279, I496
	, AL137271,
	AF061981, AF090896, AL050155, AL137550, A23630,

				S78214. D	D55641, M196	M19658, Y10080	Y10080, AL133637,
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				0	X52128, AL110221,	6110221, AI	AL050149, S83440,
				AL137660,	Y07905, A	AB029065, UE	U88966, S75997,
				AB016226,	AF100931,	AF113677,	AL117463,
				AF001215,	AL049314,	272491	
1627	HWHPZ02	876858	Preferably excluded from the	AW043824,	AI094162,	AI150332,	AW152394,
			present invention are one or more	AI363370,	AI340929,	AW341579,	AA904074,
			polynucleotides comprising a	AI015843,	AI039705,	AI192155,	AI338344,
			nucleotide sequence described by	AI038188,	AI144479,	AA922221,	AA804396,
				AA768639,	H29728, A	AA256891, AA	AA708611, H29729,
			teger between	AA902548,	AA641864,	AA256375,	AA310759,
				AL038838,	AL038983,	AA641863,	AL037727,
			15 to 1188, where both a and b	AL038532,	AI142134,	AW316536,	AA654177,
			correspond to the positions of	AL038822,	AL043814,	AL043923,	AL043845,
			residue	AL040617,	AL044186,	AL041238,	AL047012,
			NO:1627, and where b is greater	AL041577,	AL041459,	AL044064,	AL040294,
			egual to a + 14.	AL041635,	AL044037,	AL047170,	AL040463,
				AL040768,	AL046850,	AL045753,	AL041752,
			-	AL045684,	AL040625,	AL047219,	AL040052,
				AL043570,	AL043848,	AL041374,	AL043627,
				AL041523,	AL041730,	AL044074,	AL041602,
				AL043492,	AL040839,	AL043677,	AL040472,
				AL043467,	AL040510,	AL042135,	AL043538,
				AL047183,	AL040464,	AL045671,	AL046442,
				AL040621,	AL046994,	AL040444,	AL041133,
				AL039316,	AL041324,	AL046392,	AL046914,
				AL040322,	AL044258,	AL044272,	AL040119,
				AL041098,	AL041096,	AL045817,	AL040148,
				AL045920,	AL049018,	AL047057,	AL044199,
				AL044187,	AL040458,	AL041163,	AL040576,
				AL041955,	AL045990,	AL041292,	AL041358,
				AL040332,	AL041142,	AL041346,	AL040529,
				AL041159,	AL044274,		AL041168,
				AL040745,	AL046330,	AL041197,	AL040128,
				AL040571,	AL042096,	AL047036,	AL040342,

	AL044165,	0091,	AL040414,
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	AL041131,	AL039744, AL044162,	AL046327,
	AL037435,	AL040149, AL040155,	AL041051,
	AL040168,	AL044201, AL037335,	AL043775,
	AL043496,		AL037443,
	AL039432,	AL041227, AL045857,	AL040329,
	AL079878,		AA471208,
	AL041086,	AL040193, AL037323,	AW129525,
	AL040075,	AL040263, AL040370,	AL040255,
	AL038761,	AL041233, AL041140,	AL045725,
	AL039915,	AL043612, AL041246,	AL037295,
	AL041277,	AL039338, AL041278,	AL045989,
	AL049069,	AL039643, AL079852,	AL040238,
	AL043537,		AL041347,
	AL043941;	AL037341, AI028338,	AL080031,
	AL134524,	AL044125, AL037279,	AL047037,
	AL043444,	AA257137, AA629169,	AL046097,
	AA257022,	D79670, AL044529, AL045328,	.045328, AA094619,
	AL046360,	AL045994, AL042898,	AL046150, T23985,
	AL043440,	AA5854	Z30131, T19415,
	N87157, Z	Z28355, T23957, AL042712, AL038745	712, AL038745,
	AA585101,	T11028	T23888, AA585453,
	AI541374,		AI540967,
	AI546855,	_	AI541523,
	AI541514,	AI541509	AI546999, AI535639,
	AI557731,	R29445,	AI526194, AI556967, AI541508,
	R28735, A	1546945,	T41289, AI546828, AL040385,
-	AL047163,	AL079953, R29177,	AI557787, AL134110,
	AI526073,	AA585476, AA174170, AF161482,	AF161482,
	AC006530,	AR062871,	3189, A43188,
		A98420, A98423, A98432,	, A98436, A98417,
	A98427, A	A84772, A84776, A84773,	1, A84775, A84774,
	AR067731,	AR067731, AR037157, AR054109, AR067732,	AR067732, A58522,
	A91750, A	, AJ244004	767, A93963,
	A93964, A	A85395, A85476, AR062872	172, AR062873,

A25909, A81878, AF082186, A64973, A58524,	25027, I26929, I44515, I26928, I26930,	, M28262, A60212, A60209, A60210, A6	E13740, I48927, I63120, AR017907, A18050,	A23334, A75888, I70384, A60111, A23633,	2, I15717	A77095, A95051, A18053, AJ244005, I08396,	A11624,	, E01007, I13349,	A35537, A91965, A02135, A02136, A04663, A04664,	I08395, AR043601, A93016, A11245, A92133,	E12615, AR035193,	A13392, A13393, AR031488, I13521, I52048,), I49890, I44531,	E16636, I44681,	, A24782, A95117, I62368,	6, AF149828, I01995, I080	I60241, I60242, A20699, E00696, E00697, E03813,	AR009151, I66485, I66483,	I66497, I66496, AR038066,	AJ230935, AR008429, I05558	2	X07299, AJ231009, IO	D13316, AR035975, AR035977, D50010, AR009152,	AR051957, I1830	AJ238010, X81969, I19525, AR066494, M20328,	X13697, J04205, X69804, X97869, AR035974,	AR035976, AR035978, A70872, D13509, E17098,	X14684, AJ231028, I66495, I66494, A22734,	AR022273, X91336, AJ230867, AJ230845, A70869,	I36244, A29109, A32111, AR051864, D17247,	A93923, AR051865, A06631, S60422, A83642,
																									-						

				166491, 166492, 166493, 166481, A83151, A93916
				2 AROSREK4 AS4E4R AS4E4E V14219
			-	, IO5845, X91337, AC005541, AA97181
				AI032717
1628	HLTAZ90	876865	Preferably excluded from the	AA873435, AA600839, AI768313, AI146480,
				, AA773760,
			ot Ct	AI936013, AI887319, AW247888, AI290267,
			e sednence	AI949176, AI140850, AI383970, AA478888,
			al formula of a-b, where	AI335758, AA455467, AI131375, AA446062,
			is any integer between 1 to 1375 of	AI375904, AW273478, AI569525, W92189, AI080606,
			SEQ ID NO:1628, b is an integer of	AA446800, AI922678, W48604, AI669705, AI088017,
		_		AI079611, AI357729, W94886, AA778027, AI420677,
			correspond to the positions of	AA662489, AA199802, AA199694, N99008, AA455466,
			le residue	.097343, RE
			NO:1628, and where b is greater	AW079086, W81498, AA478769, AA602304, AA770587,
			than or equal to a + 14.	
•		_		AI687665, AI275990, AI127693, AI040179, H06586,
				AI188614, AI383744, AI160662, T16066, AW162694,
				_
				, AA577605,
				_
				N75810, AA999862, AA417649, AA582611, AI400342,
-				AA749354, AA923020, AI537750, AI579976,
				, AI915035, N69819, AA
				, AI433790,
				6, AI207126, AA470409,
				AA074998, AI432068, AA725585, AA757124, N75636,
				R50657, H06531, T27805, AI220764, W81497,
				AA492209,
				AI524835, T59434, AA035575, AA364008, D12231,
				, AA482915
				348,
				AI932950, R29196, D12095, AA343259, AA588441,

				T09050. AI239988, AI572155, T33940, AI917677,
				3, AI915005, D57719, AA490946
				, AA659260, R15055,
				AA534940, AW262956
				AA902888, AA736627, T09051, AA491132, AI557731,
				AC004081, AC007666, AC000052, AC004019,
·				AF055664, L08069, D13388, U53922, AA446079,
				AA429922
1629 HHF	HHFUM32	876866	Preferably excluded from the	AA525015, AI097213, AI186110, AI205864,
			present invention are one or more	AI460279, AA454512, AW003859, AI143331,
			polynucleotides comprising a	AI305240, AI337532, AI279156, AI333362,
			nucleotide sequence described by	AA770652, AA483013, AA846308, AI024319,
			the general formula of a-b, where a	AI380066, AI184498, AI204185, AI332737,
			is any integer between 1 to 607 of	AI025452, AA701068, AW298191, AA314391,
			SEQ ID NO:1629, b is an integer of	AA780879, AI204046, AA722950, AA903838,
			15 to 621, where both a and b	AI368078, AI073640, AA010086, AA911716,
				AA948332, AI188877, N45102, AI094300, W52409,
			Φ	AI311092, AA622052, AI302571, AI369905,
			NO:1629, and where b is greater	AI660241, AI138619, H48026, H41034, AI749308,
			equal to a + 14.	N76689, AI354731, N31297, AI141562, AI347212,
				A1191310, A1092132, AA875920, AI346333,
				AI344362, AI186141, AI184174, N50933, AA854247,
	_			W32499, H93326, AA740175, AA765339, AA886065,
				AI718470, N54609, F32533, AA229525, AA604454,
				R97891,
				AI312692, N46264, AI027037, AI192124, W77745,
				AA629102, AA975984, W05153, N45023, R68274,
				AA046489, AA362565, W99330, AA075564, H18704,
				œ
				AA887933, H41035, H23703, N84980, N69892,
				AA311757, H18805, F36632, R26083, AA046701,
				AI702033, H18369, AA327843, AA299086, F33066,
				R68309, W52410, AA877022, AA643367, AA079015,
				5, H26911, H57271,
				R96486, AA339947, W02163, AI220631, W05365,

				AA772749, H28518, H A46263, A	, H93830, F26046, H23704, AA083351, AA35275, AA02476		H58286, R94598, AA075559, AA296237, F33965, AT557901
					, w w	F28514, AI750084, R47744, AW265596.	
				R50391, A		A659764, A	AA659764, AA302180, W31292,
				N	C00512, A	AA709422, F	524, AL08
						X69904, X05218,	D13123,
				M16453, T	T80797, T81	T81201, H27411, N93425, N95193	1, R97890, N41011 3 W24594
				ဖ	v		
1630	HHFAB62	876870	bly exclude	AA346386,	AW300186,		AW364745,
			present invention are one or more	AW374001,	AW364749,	AW373998,	AL046035,
			polynucleotides comprising a	AW373994,	AW364756,	AW373996,	AW373989, D79991
			nucleotide sequence described by				
			the general formula of a-b, where a				
			is any integer between 1 to 1144 of				
			SEQ ID NO:1630, b is an integer of				
			15 to 1158, where both a and b				
			correspond to the positions of				
			NO:1630, and where b is greater				
			than or equal to a + 14.				
1631	HWLWJ70	876873	Д	AA527360,	AW051577,		AI590246,
			present invention are one or more	AA482382,	AA417897,	AA834979,	T33217, AI933007
			polynucleotides comprising a	AA886393,	AI242582,	AI242582, AA912932,	AA552566,
			nucleotide sequence described by	AA026889,	H12586, A	H12586, AA770351, AI122821,	1122821, 245211,
			wher	AA810545,	AA089741,	AA026890,	AW235276,
				AA442516,	AI081311		
			SEQ ID NO:1631, b is an integer of				
			15 to 679, where both a and b				
			correspond to the positions of				
			nucleotide residues shown in SEQ ID	****			
			than or equal to a + 14.				
1632	HCRPV85	876876	Preferably excluded from the	AI138310,	AA579608,	AL080041,	AA150112,

		present invention are one or more	-	AW139942, AI669978,
		polynucleotides comprising a	AA150453, N21199, AW	AW337765, W60839, AA007492,
				H39087, AI857928, AW402945,
		~		AW044377, AA708593, H06461,
		ny integer between 1 to	AI554400, AA806848,	AA292984, AA281307, D53188,
		SEQ ID NO:1632, b is an integer of	AI074110, AI359733,	H37969, AA911725, AA194095,
		, where both	AA757126, AA815284,	AW166409, AI362093,
		correspond to the positions of	AA258691, AW386068,	AA614128, AI937918,
		residue	AI218676, AA429422,	AI361580, AA156587,
		NO:1632, and where b is greater	AA931474, N27470, AA	
		equal to a + 14.		D52529, AA171394, AW367949,
	· <u>-</u>		AA150166, W47135, AA	D5
-			AA937690, AI752560,	AA312520, AI039854,
			AI282901, AA884648,	AI094728, AI201298,
			AI273365, AI346383,	AI421258, AI310120,
			AI361451, AI285056,	AA040411, AA789206, N88385,
			AI418521, AI973164,	AA227133, N99005, AL038896,
	_		AW362878, AW403348,	W24127, AL119637, AI016520,
			AA541481, AA309620,	AA150397, AA306805,
			AI400189, AA284235,	H51237, AA331743, AW023315,
			R67309, AA373361, AA156654,	156654, AA730527, D57421,
-				W69686, AA332449, AI368439,
			AA281258, AA040303,	AA040303, H63313, AA359717, AW362873,
			R74438, AA770542, H06565,	16565, AA363548, AI339537,
			7, AA884006	, AA922473,
			A	AA722328, AW207758, AI753879,
				AA999738,
			AW151651, D52528, AW	AW391062, AA330258, Z45721,
				AA484242, AA382542,
-				
				D53164, AW386086, AI749497,
•			3, H68127,	AI149688, AA365933, H11545,
			R26679, R36441, AA70	
			T30044, T85823, AA35	73, R57470
			Z42383, D53068, N71806	106, D53095, C03662,

				AA263144, AA111835 AIO56722, N21394, N63858, AA503313, AA290658, R60888, AL037148, X74262, AE000658, U85195, AA703653, AA853715		, D81554, T83223, AA354104, AW270594 F23396, AI973191, AA382087, AA677495 X71810, U35141, AF AC005277, AA045875	AA354104, AW270594, AI571557, F23396, AI973191, AI590666, AA382087, AA677495, AA290659, X71810, U35141, AF097750, AC005277, AA045875, AA398311,
1633	HCE3V58	876878	inv inv sot ral ral ral ral ral an	AW301835, AA878213, AI285166, F35821, H	, AI308020, , AA694197, , AA133903, H90906	AI860966, AA088689, AA302740,	AA133904, AA133904, F26419, AA582580,
1634	нкавен	876882	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 3629 of SEQ ID NO:1634, b is an integer of 15 to 3643, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1634, and where b is greater than or equal to a + 14.	AIS24051, AI401617, AW021551, AA856594, AW07900, AI191795, AA1999, AA579953, AA579953, AA579953, AA579953, AA579953, AA1093279, AI306480, AI306480, AI306480,	AMO07724, AI936772, AI923848, AI802150, AI680189, AI160059, AI128436, H99041, A AA702093, AA702093, AA71204219, AI123115, AI123115, AI123115,	A1609303, A1735659, AA889466, A1393945, AN246695, AN246695, AN274108, A1274108, A1015928, AZ AA453200, A1026873, A1026873, A1075685, A1373875, A1373875, A1373875, A1373875, A1373875,	3, AI560001, 9, AI249001, 6, AI770052, 5, AI963008, 1, AW081918, 6, AA812522, 8, AA909840, AA931655, AI262534, 7, AI359209, 9, AI125919, 9, AI125919, 9, AI183377, 9, AA169284,

	AI690264, AI356799, AI298090, AA847328,	
	3, AI573004,	
	5, AA398429, AA781758, AI24861	
	AA972778, AL120931, AA813433, AW364708,	
	AI625940, AA975860, AI051123, H18709, AI624093,	093,
	I299217, AI149399,	712,
	N90938, AW373562, AA687	849,
		703,
		641,
	, AW131426	5,
	W04345, AI348671, W17386, AA804381, AA305682,	2,
	I, AA291227	526,
	, AA732854,	
	AI080704, W68454, AI084772, AW084472, AA479223,	223,
	N27564, AI184963, AA505251, AI022978, D53877,	7,
	A	734,
	AI245609, AA454161, AI589126, AI690281, N92279,	279,
	AA745905, AW057830, AA467899, AA938231,	
•	7.	
	AI890992, AA847408,	1,
	AA456530, AA971614,	
	AA614659, AA788753,	-
	AA454984, AW168929,	
	, AA151101	7
	I298694, W17235, AI348194,	
	, H18598, N89744,	-4,
	A535636, D6	,6
	AI433476, AI248821, AI245176, AA403052,	
	, AA150090, W17187, W16652, AA	7,
	AA031722, AA089961, AI287545, T75133, R75976,	, 6,
	, AI245198, AI814870,	851,
	S, AA701210, AA578436, AA15	
	67843, D53769, N22090, D52639, T28569,	
	AA889910, D56802, AI890986, C01613, H88627,	

			AI267864, H46858, AI075972, AI242237, AI185100,
			, H88628, AA968979, Wl
			, ноз189,
	-		AC004520, M29065, M29064, AF073993, AF192348,
			09123, L02954, L0295
			9, R12
			887, R23576,
-			R52479,
			H03988, H04178, N34507, N40385, N74179, N74343,
			N93493,
			W17019,
			W81443, N90151, AA036938, AA167390, AA483158,
			AA632646, AA765452, AA808476, AA888709,
			C02214
			AA092059, AA211492, AA216333, Z20376, AA703571,
		•	AA889282, AI032462,
			D20533, T24609, F013
1635 HRAEG13	876886	Preferably excluded from the	, AL079428, AI962210,
		present invention are one or more	AW409972, AW362305, AW410672, AI924517,
		polynucleotides comprising a	, AW025356, AA405914,
		nucleotide sequence described by	, AI523918,
		the general formula of a-b, where a	AW206660, AI569743, N94878, N99556, AW301065,
		ě	354, AI936512,
		SEQ ID NO:1635, b is an integer of	ω,
		both a and	2,
		positions c	I347352, AW387060, AW386988,
		de residues s	AI081389, AA350220, AI148131, AA783037,
		NO:1635, and where b is greater	796, AI277386,
		than or equal to a + 14.	AI372627, AA4053
			ر و
			AA781626
			R56864, R55500, T66335, H92624, AA350276,
-			R81346, AL121276, AA350037, F09706, AI298408,
			79, R51360
			F12065, H50110, AA351242, N22306, F09146,

AA234354, N26102, N55429, AL120770, AW387043, AA405389, H50154, H43762, AW387110, H72992, AA227365, R79738, R79737, H44600, H70095, R50621, AI184049, R45951, H29909, T66284, AA744978, N71548, H72991, AA368705, AA936885, AI739624, R55499, AW007986, T83200, AI863755, R50454, R50527, T36310, R50455, T85587, T77076, AA936368, H43432, AA464051, T87308, T07160, T78532, AA321966, AW268156, T85586, H43431, F26601, N40316, AI832126, AI372626, AW376436, N54476, R81601, R51465, R94300, AW367002, AA324819, N76802, AW073570, AI654772, AI473579, N76587, F35806, H92406, AW366992, AA302603	, , ,	AA946784, AW375919, AA527581, AA904758, AA209387, AA563949, AI833239, AA740268, AA527668, AW372169, AA948567, AA894539, AI745625, AA468774, AA725505, AW376020, AA164354, AA946619, AI348033, AA594622, AA53342, AW160477, AA937588, AA662503, AA588618, AW363501, AW375476, AA677897, AI310309, AI123763, H59915, AW161438, AW160982, AN160317, AI907434, AA780152, AW363508, AA526226, AW295010, AW176047, AI472327, T65562, AI005477, AA349978, AA928712, T08552, AA610643, AA349672, T65630, F12026, AA434132, AW365033, AA349672, T65630, F12026, AA434132, AW365033, AA38674, AA453217, AA384272, AA339261, AA3867135, T03912, R78158, AA367413, AA357314, H27129, R91610, AI766762, D51350, AA343589, T03852, AA384370, AW264753, AW376759, AW376799,
		Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 1228 of SEQ ID NO:1636, b is an integer of 15 to 1242, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1636, and where b is greater than or equal to a + 14.
		876888
		HLIB207
		1636

				AW376653, AA362098, D54438, AI905702, AA300134,
				H2
				, S75311, A
				되
1637	HTPFB46	876890	Ω	AW444886, AI983059, AL135147
			present invention are one or more	W07327, AI492267, AI
			polynucleotides comprising a	, AA406085, AI678761,
_			nucleotide sequence described by	
			the general formula of a-b, where a	AA804950, AA533437, AI242554, AI223449,
			is any integer between 1 to 2110 of	, AA644395,
			SEQ ID NO:1637, b is an integer of	
			15 to 2124, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	AA932178, R78020, AI089059, R32384, AI242914,
			NO:1637, and where b is greater	T81104, F34121, AI468126, F25882, N75820,
			than or equal to a + 14.	AI335792, F35752, F18999, AI984724, AW305237,
				AW086410, AW272065, AI310836, AI345115,
				AI223675, AI308339, AI312490, AI252159,
				AI307405,
				AI252373, AI349681, AI252335, AI250483,
				AI252345, AI583501, AI583500, AW302935,
				AI583889, AW303168, AI348995, AI349742,
				AI309420, AW269095, AI336494, AI335439,
				AI349287, AI306795, AW274358, AI349945,
				φ
				L40817, L44140, X87196, X74606, X90393
1638	HDPSS23	876892	Preferably excluded from the	AI129800, AW027959, AI927949, H92980, AI650270,
			present invention are one or more	AI708393, AL138076, AA524072, AI831594,
			polynucleotides comprising a	AA749139, AI926721, AI399955, AI302816,
			nucleotide sequence described by	AA262795, AI862160, AI093249, AA828301,
			the general formula of a-b, where a	AI625105, AA904444, AA772552, AI816834,
			is any integer between 1 to 1421 of	AI084565, AA314418, N30447, AI242763, AI810709,
			SEQ ID NO:1638, b is an integer of	AI653617, AI129801, AA443839, AI289975,
			15 to 1435, where both a and b	AA281653, N25206, AI758575, AA026905, AA737455,

	correspond to the positions of	AA039864, AW000990
	nucleotide residues shown in SEQ ID	AI032004,
	NO:1638, and where b is greater	AW084297, R97735, AI640264, AA336497, AW080103,
	equal to a + 14.	
	•	C21440, AA3387
	-	N69415, N91446, R764
		, AA281785, AA680378, T185
		T10789, AA610255, AA568204, AA570740, AA483606,
		T47138, AW151018, AI355246, AI445373, AI915081,
		AA219349, AA664126, AA582746, AW275432,
		AA558404, AA837771, AA214453, AA857812, T94394,
		AA482792, AI249688, AI567391, AA630854,
		AA683069, R67701, AA515939, AA425924, R77139,
		AW069227, AA714073, AA297006, AI285493,
		AI298079, R79929, F35097, AI634377, AI791659,
		AW104163, AI671077, AL048060, AA809186,
		AA831408, F35684, AW084967, AA523695, AI962030,
		AI185394, AA491767, N51636, AI538236, AA558366,
		1, AI735092, AA3763
		•
		A
		AW419389, AA632556, AI634187, AA302978,
		AI457313, AI620992, AI358542, AA769141,
		AA342238, AA583386, AI312090, AI049630, U91323,
		AC004686, AL080245, AL035587, AC002073, Z81357,
		AP000045,
		AC005702, Z82901, AC007774, AP000030, AL008718,
		AC004079,
		AC005759, AC002365, AC007193, Y07848, AC004598,
		AL096701, AC002565, AC005799, Z73900, AC007390,
٠		AL031721, 293016, AL118497, AC006501, AC007566,
		AC000064, AP000133, AP000211, AC004859,
		AC006333, AC007179, AC000025, AL049776,

Т	· .	
AL078644, Z94802, AF064861, AC006121, Z98051, AL049610, AF102137, AL008582, AP000555, AC009247, AL049843, AC007899, AC004974, AC007172, AC06120, AC008149, AC004780, AP000355, AL049643, U78027, AC006276, AL035450, AC005089, Z93784, AC005399, AC006430, AC007114, AC002550, AC004587, AL022316, AA261881	AI676066, AA87 AI862396, AW134 AI760422, AW34 AI950211, AI36 AI850709, AI52 AA1860709, AI52 AA319888, AA942 AA319888, AA0842 AA319888, AA0842 AA319888, AA0842 AA319888, AA084 AA5251, W72425, AA011178, AA03 AA044195, AA08 AA038458, AA94 AA477226, AA70 AA661647, AA80 AA661647, AA80 AA63245, AA09 AA132001, AA18 AA432945, AA09 AA132046, AA694 AA432945, AA02 AA132949, AI120 AI139707, AI20	AA629925, AI557066, H72652
	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 1617 of SEQ ID NO:1639, b is an integer of 15 to 1631, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1639, and where b is greater than or equal to a + 14.	Preferably excluded from the present invention are one or more polynucleotides comprising a
	876901	876903
	HCEIC29	HE90Y91
	1639	1640

			nucleotide sequence described by the general formula of a-b. where a	
			ய	
			SEQ ID NO:1640, b is an integer of	
			15 to 853, where both a and b	
	,		correspond to the positions of	
			ъ	
			NO:1640, and where b is greater	
			than or equal to a + 14.	
1641	HFKFN66	876904	Preferably excluded from the	AL031433
			present invention are one or more	
			polynucleotides comprising a	
			nucleotide sequence described by	
			the general formula of a-b, where a	
			SEQ ID NO:1641, b is an integer of	
			15 to 688, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1641, and where b is greater	
			than or equal to a + 14.	
1642	HWMFQ16	876905	\sim	AA775776, AI041206, AI884423, AA608631,
			present invention are one or more	AA307942, AA602534, AA477709, AA604331,
			polynucleotides comprising a	AA610041, AA237053, AI874354, AI922651,
	_		nucleotide sequence described by	AA455372, AA478920, AI861817, AI174744,
			al formula of a-b, where	, AI803985, AA307739,
			teger between 1 to 190	8, AI420956
			SEQ ID NO:1642, b is an integer of	ω,
			15 to 1916, where both a and b	AI369854, AW402584, AA250883, AI362747,
			correspond to the positions of	_
			യ	r Ž
			NO:1642, and where b is greater	, AA20654
			than or equal to a + 14.	AA908393,
				AA205036, W07733,
				AA723847, AA151196, R68884, AI217962, N62289,
				R60986, AA019523, AI307617, AA535112, H18659,

HCRBB01 8769	4 9	AA484614, AA252156, AA394239, N89897, AA418016, AI289322, N35239, T96813, T98004, Z39105, AI347692, AA401922, R68786, AI421701, AA300711, AI984054, AI307367, AI869880, AW003896, AI357580, AI097540, H78257, AA773528, AI933853, N26474, W19451, T96826, AA937255, AA494127, AA456012, AA622190, AA531018, AW264334, AA296375, AW340846, R39778, AW368305, T98082, AW406763, AW389979, T32639, AW389990, AA773673, AA304962, AA233500, AW383537, R58298, C15957, N78713, AA019294, D78788, AW389995, AA402093, D31588, AW366573, AA095078, N87188, N86592, N88113, N88337, N85682, AF078859, AF078868, AL021878, AF090946, U21721, AJ243486	Preferably excluded from the Present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a sequence described by the general formula of a-b, where a sequence described by the general formula of a-b, where a sequence described by A1749224, AA307941, AW272874 AN274596, AI336326, AW169351 SEQ ID NO:1643, b is an integer of AV274596, AI336326, AW169351 AV269482, AI749219, AI026046 AIS910185, AW361012 AI922602, W60954, AI735165, AI931683, AI931689, AIS9188, AIS88266 than or equal to a + 14. AI653978, AI890155, AI934802
			876909

A118 A118 A118 A118 A419 A4118	7, AA555069, AA860461, AI68937 10, AI937827, AI003581, AI83136 10, AM377974, AI084421, AI09208 11, AA505597, AI336527, AA55468 12, AA605597, AI336527, AA55468 13, AA798382, AW083700, AI03873 14, AI073882, AI031884, AA82638 17, AI625287, AI031884, AA82638 17, AI625287, AI022580, AW27058 18, AI073697, AI032680, AA75908 19, AI073697, AI354660, AA52612 19, AM150128, AA638242, AA55442 19, AM150128, AA60805, AI96310 19, AA7235239, AW089108, AA50822 20, AI572298, AA382418, AA50822 21, AIS88932, AI918522, AA91605 22, AI572298, AA382418, AA5982 23, AA7235239, AW089108, AI33527, AA29959 24, AIS88463, AA216394, AA59829 25, AI572298, AA382418, AA34115 26, AA837555, AI933527, AA29959 27, AA582932, AI933527, AA59959 28, AA650299, AA56817, T27681, AS9, AA550299, AA550294, AI72087 29, AA650299, D56517, T27681, AS9, AA550299, AA555204, AI72087
AWI.7	AWI/6624, AW36/125, AA146683, AI161032,
AA220	AA226022, H88875, H88876, AA523823, AA302252
AA300	AA301829, W21502, W70311, AA311904, AA720066

				AI273789, AA096200, AW377515, AA729962, D20952, AI811103, T84076, X60111, AR016441, I13744, M38690, D10726, AC006057, L35275, M81720, L08115, D30786, AR016440, E05732, X76489, L08125, L08118, U15792, S60490, L08119, L08120, L08122, L08123, L08124, L08121, S60489, S60462
1644	HSAAN15	876912	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 1095 of SEQ ID NO:1644, b is an integer of 15 to 1109, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1644, and where b is greater than or equal to a + 14.	AA643028, AI858075, AA910344, AA573333, AI088151, AA481497, AA826812, H63145, H0 AA521057, R53520, AA AA732248, AA970100, R53519, AA373512, R4 A481183, AW207413, AI
1645	HTEKS27	876913	preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 2159 of SEQ ID NO:1645, b is an integer of 15 to 2173, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1645, and where b is greater than or equal to a + 14.	AI657156, AI375103, AI684065, AA678409, I655208, AI702778, AI R54239, AA436083, R5 A112078, R35463, L138 59697, R51845, AI4792 AI635429, L13826, R3 61243, L23208, AR0513 23311, AR051321, L301
1646	HWMBA1 0	876920	preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 1380 of	AI749171, AI660550, AA677676, AA464420, AA284905, AA718994, AI141193, AA481894, AI078424, AA481977, AA703408, AI276556, AI017050, AA502348, AA936362, AA936704, AW131471, F36806, AW273475, AI261777, AI218960, AI218966, AI744229, AI248232, AA452839,

			SEQ ID NO:1646, b is an integer of	AI277984,	AA053718,	AI150864,	AI140517,
				AI129769,	AI160406,	AW152129,	AW000750,
				AI248566,	AI805790,	AI826304,	
		_	nucleotide residues shown in SEQ ID	AA020812,			AA019875,
		·-·.	-	AW242786,			F22534, AI240050, T41072,
			than or equal to a + 14.	W96529, AW069782,		68326, AAO	W68326, AA053858, H37782,
				AA055112,		AI765563, F	F31495, AA020811,
				AI244397,	H37923, A	A013192,	T51835, R50369,
		4		AW339481,	AI903705,		AW194148, AA019902, W68142,
				AW298469,	AW003689,		AA019913,
				AW139654,	AA383551,		AA384419, AA883222, H41086,
							H86062, AI735754, R80952,
				W92479, A	AA535061, F	31376, T40	F31376, T40204, C04332,
				AA019941,		AW050973, AI560455	AI560455,
				AI470969,		I695746, A	AI695746, AA284774, AA855078,
		-		AA013427,	H38276, W	92489, AA4	W92489, AA412431, AA844626,
				AW074589,	ဖ	H86397, AA906632,	A906632, F36956,
				AA018714,	AA021006,		
				AA015696,	AW050422,	AA402869,	AA015660,
				AA464421,	AA454730,	AA015659,	AA454780, T28267,
				AA018985,	AA018750,	AC006449	
1647	нсовозя	876921	Preferably excluded from the	AI803478,	AA578800,	AI760557,	AA569728,
			inventic	AI803206,	AI199737,	AI524625,	AA825640,
			polynucleotides comprising a	AA937979,	AI436327,	H83996, A	H83996, AA879427, AW205011,
			nucleotide sequence described by	AI284171,	AA262130		
			the general formula of a-b, where a				
			is any integer between 1 to 711 of				
			SEQ ID NO:1647, b is an integer of				
			15 to 725, where both a and b				
			correspond to the positions of				
			nucleotide residues shown in SEQ ID				
			NO:1647, and where b is greater				
			than or equal to a + 14.				
1648	HWLGQ64	876923	Q	AI743526,	AA535976,	AA534299,	AI245191,
			present invention are one or more	AA917952,	AI360198,	AA189088,	AI476640,
			polynucleotides comprising a	AI750101,	AI151214,	AI219288,	AI189990,

				4480 700 4 80
1649	HCQCV14	876926	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 558 of SEQ ID NO:1649, b is an integer of 15 to 572, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1649, and where b is greater than or equal to a + 14.	AP000529, AP000528
1650	HCROO59	876934	preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 391 of SEQ ID NO:1650, b is an integer of	AA376902

			15 to 405, where both a and b					
			nd to the positions of					
			nucleotide residues shown in SEQ ID	_				
			NO:1650, and where b is greater					
			than or equal to a + 14.					
1651	HCRPN27	876936	Ω	AA457220,	AA354909,	AA040828,	AI688798	
			present invention are one or more					
			polynucleotides comprising a					
			nucleotide sequence described by					
			the general formula of a-b, where a					-
			is any integer between 1 to 981 of					
			SEQ ID NO:1651, b is an integer of					
			15 to 995, where both a and b					
			correspond to the positions of					
			nucleotide residues shown in SEQ ID					
			NO:1651, and where b is greater					
			than or equal to a + 14.					
1652	HCRON34	876938	Preferably excluded from the	AI634562,	AA129701,	AA129323,	AA129745,	
			present invention are one or more	AI269483,	AI952719,	AI656261,	AI239764,	
			polynucleotides comprising a	AI678885,	AI873730,	N48153, A	AI873730, N48153, AA904475, AA65351	653518,
			nucleotide sequence described by	AI538894,	R43961, AI287295	I287295, W	W68609, AI114476	4476,
			the general formula of a-b, where a	AA973355,	AI866872,	AA133249,	AI681503,	
			is any integer between 1 to 622 of	AA133292,	AI690203,	AW271391,	D29021, AI	AI186074,
			SEQ ID NO:1652, b is an integer of	AA757303,	AA742226,	AA73777,		AI825401,
			15 to 636, where both a and b	AI934240,	AA587412,	AW051055,	AW020046, W68807	W68807,
			correspond to the positions of	D83781				
			nucleotide residues shown in SEQ ID					
			NO:1652, and where b is greater					
			than or equal to a + 14.					
1653	HFKFH50	876940	Preferably excluded from the	AA927698,	AI300925,	AW009795,	AA402380,	
			present invention are one or more	AI830852,	AA430318,	AI493302,	AI142868,	
			polynucleotides comprising a	AI037989,	AI423267,	W52884, A	AA907276, AI333045	333045,
			nucleotide sequence described by	AA628712,	AA988209,	AI363130,	AA987992,	
			the general formula of a-b, where a	AA578507,	AI298580,	AA639466,	AA402235,	
			is any integer between 1 to 1241 of	AI052201,	AI073629,	AA458463,	AA564499, 1	N78968,
			SEQ ID NO:1653, b is an integer of	AA534799,	AW083734,	AA442975,	AI074925,	

EQ ID AA425157, r R37964, W3 R37964, W3 R446682, AA485845, AA031677, AA053511, AI867052, L05489, M3 X67295, L1 X67295, L1 X67295, L1 AW392670, AW363220, AW363220, AW119336, AL119336, AL119336, AL119336, AL119336, AL119496, AL119496, AL119496, AL119496, AL119496, AL119496, AL119496, AL119496, AL119496, AL134533, AL037054, AL037526, AL037526, AL037054, AL037526, AL036889, AL037656, AL037526, AL037526, AL037656, AL036894, AL036894, AL0366436, AL0366436, AB026436, AB026436,			15 to 1255, where both a and b	AA053124, C04884, AA775515
nucleotide residues shown in SEQ ID AA359764, WO:1653, and where b is greater R37964, W3 WO:1653, and where b is greater AA46682, AA031677, AA031670, Desent invention are one or more AM302670, present invention are one or more AM302670, present invention are one or more AM302670, the general formula of a-b, where a AL119391, is any integer between 1 to 504 of AL119395, SEQ ID NO:1654, b is an integer of AL119395, IS to 518, where both a and b AL119395, IS to 518, where both a and b AL119395, IS to 518, where both a and b AL119395, IS to 518, where both a and b AL119395, IS to 518, where both a and b AL119395, IS to 518, where both a and b AL119395, IS to 518, where both a and b AL119395, IS to 518, where both a and b AL119395, IS to 518, where both a and b AL1194957, than or equal to a + 14. AL119496, AL0310954, AL032044, AL0310056, AL032043, A			to the positions of	AA425157, R83528, AA401316, AA676435, D51268,
NO:1653, and where b is greater than or equal to a + 14. HCRQG66 876941 Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 504 of SEQ ID NO:1654, b is an integer of 15 to 518, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1654, and where b is greater than or equal to a + 14. HCROW80 876942 Preferably excluded from the polynucleotides comprising a nucleotide sequence described by			residues shown in SEQ	AA359764, H27189, C01185, AA402234, H27190,
HCRQG66 876941 Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 504 of SEQ ID NO:1654, b is an integer of 15 to 518, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1654, and where b is greater than or equal to a + 14. HCROW80 876942 Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by			and where b is	39595, T27801, D55114, R45640
HCRQG66 876941 Preferably excluded from the present invention are one or more polynuclectides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 504 of SEQ ID NO:1654, b is an integer of 15 to 518, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1654, and where b is greater than or equal to a + 14. HCROW80 876942 Preferably excluded from the polynucleotides comprising a nucleotide sequence described by			equal to a +	
HCRQG66 876941 Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 504 of SEQ ID NO:1654, b is an integer of 15 to 518, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1654, and where b is greater than or equal to a + 14. HCROW80 876942 Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by				_
HCRQG66 876941 Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 504 of SEQ ID NO:1654, b is an integer of 15 to 518, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1654, and where b is greater than or equal to a + 14. HCROW80 876942 Preferably excluded from the polynucleotides comprising a nucleotide sequence described by				
HCRQG66 876941 Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 504 of SEQ ID W0:1654, bis an integer of 15 to 518, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1654, and where b is greater than or equal to a + 14. HCROW80 876942 Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by	-			AA343828,
HCRQG66 876941 Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 504 of SEQ ID NO:1654, b is an integer of 15 to 518, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1654, and where b is greater than or equal to a + 14. HCROW80 876942 Preferably excluded from the polynucleotides comprising a nucleotide sequence described by				AI867052, AC004634, AR042382, L17032, L36027,
HCRQG66 876941 preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 504 of SEQ ID NO:1654, b is an integer of 15 to 518, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1654, and where b is greater than or equal to a + 14. HCROW80 876942 Preferably excluded from the polynucleotides comprising a nucleotide sequence described by				3012, X89728, Y15731
HCRQG66 876941 Preferably excluded from the present invention are one or more Aw363220, Ab119484, polynucleotides comprising a https://doi.org/10.1001				X67295, L17029, L17030
present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a lis any integer between 1 to 504 of SEQ ID NO:1654, b is an integer of nucleotide residues shown in SEQ ID nucleotide residues shown in SEQ ID NO:1654, and where b is greater NO:1654, and where b is greater than or equal to a + 14. HCROW80 876942 Preferably excluded from the polynucleotides comprising a nucleotide sequence described by ALO330056, ALO33650, ALO33650, ALO37034, ALO38659, ALO37034, ALO38650, ALO37034, ARO38650, ARO37034, ARO38650, ARO37034, ARO38650, ARO37034, ARO38650, ARO37034, ARO38650, ARO37034, ARO38650, ARO37034, ARO37034, ARO37034, ARO38650, ARO37034, ARO	╁	-	excluded from	l
polynucleotides comprising a hill9497, ALI19444, nucleotide sequence described by the general formula of a-b, where a ALI19319, ALI19324, the general formula of a-b, where a ALI1931, ALI19522, is any integer between 1 to 504 of ALI19395, ALI19418, SEQ ID NO:1654, b is an integer of ALI19396, ALI19341, alight correspond to the positions of ALI19496, ALI19496, ALI19496, ALI19496, ALI19496, ALI19496, ALI19499, ALI19496, ALI19499, ALI194999,			present invention are one or more	, AL119484,
the general formula of a-b, where a AL119319, AL119324, the general formula of a-b, where a AL119391, AL119522, is any integer between 1 to 504 of AL119395, AL119318, SEQ ID NO:1654, b is an integer of AL119396, AL119341, 15 to 518, where both a and b AL037051, AL043147, correspond to the positions of AL119496, AL134524, NO:1654, and where b is greater AL134527, AL134528, than or equal to a + 14. NO:1654, and where b is greater AL036858, AL036859, AL036859, AL037094, AL038509, AL042544, AL037094, AL038509, AL042544, AL037094, AL037094, AL037094, AL037094, AL037094, AL037095, AL037094, AL037095, AL0			polynucleotides comprising a	
the general formula of a-b, where a AL119391, AL119522, is any integer between 1 to 504 of AL119335, AL119418, SEQ ID NO:1654, b is an integer of AL119366, AL119341, 15 to 518, where both a and b AL19496, AL134510, nucleotide residues shown in SEQ ID AL036858, AL036824, NO:1654, and where b is greater AL134527, AL134528, than or equal to a + 14. HCROW80 876942 Preferably excluded from the PA330056, AA236014, AR036034, AR0			nucleotide sequence described by	
is any integer between 1 to 504 of SEQ ID No:1654, b is an integer of ALI19336, ALI19418, 15 to 518, where both a and b ALI1946, ALI19451, correspond to the positions of ALI19496, ALI194530, nucleotide residues shown in SEQ ID ALI34531, NO:1654, and where b is greater ALI19457, ALI194599, than or equal to a + 14. ALI19457, ALI19499, ALI3939, ALO4254, ALO38509, ALO37094, ALO38509, ALO37094, ALO38509, ALO37094, ALO38520, ALO37094, ALO3809, ALO			l formula of a-b, where	_
SEQ ID NO:1654, b is an integer of a L037051, AL043147, AL035725 15 to 518, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID AL036858, AL036924, AL134531 NO:1654, and where b is greater than or equal to a + 14.			eger between 1 to 504	, AL119418,
15 to 518, where both a and b aL037051, AL043147, AL036725 correspond to the positions of nucleotide residues shown in SEQ ID NO:1654, and where b is greater than or equal to a + 14. NO:1654, and where b is greater aL134533, AL119399, AL042984, AL042989, AL042551 aL042551, AL042989, AL042989, AL042989, AL042989, AL043018 aL04264, AL037084, AL037085, AL0			1654, b is an integer	, AL119341,
No.1654, and where b is greater AL119496, AL134530, AL134515			15 to 518, where both a and b	_
NO:1654, and where b is greater			_	
NO:1654, and where b is greater AL134527, AL134528, U46346, than or equal to a + 14.			residues shown in SEQ	, AL036924, AL134531
than or equal to a + 14. AL134533, AL119399, AL042975, AL042542, AL042544, AL042989, AL037094, AL038509, AL037526, AL037085, AL037526, AL037677, AL119464, AL038520, AL036733, AL037027, AR066494, AR060234, AR066494, AR060234, AR066494, AR060234, AR066494, AR060234, AR066494, AR060234, AR066494, AR060214, AR066494,			NO:1654, and where b is greater	, AL134528,
AL042975, AL042542, AL042543, AL042543, AL042989, AL037094, AL038509, AL037094, AL038509, AL037094, AL037094, AL038509, AL037526, AL037085, AL037526, AL037087, AL036733, AL037027, AR066494, AR0602344, AR066494, AR0602344, AR066494, AR0602344, AR06494, AR0602344, AR06494, AR0602344, AR06494, AR0602344, AR06494, AR0602344, AR06494, AR0602344, AR06494, AR06494, AR0602344, AR06494, AR06026436, AR06494, AR06			than or equal to a + 14.	_
AL042544, AL042989, AL037094, AL038509, AL037094, AL038509, AL037094, AL038509, AL037526, AL037085, AL037526, AL03767, AL119464, AL038520, AL036733, AL037027, AR066494, AR060234, AR066494, AR060234, AR066494, AR060234, AR06494, AR06494, AR060234, AR06494, AR06494, AR06494, AR060234, AR06494, AR06494				_
HCROW80 876942 Preferably excluded from the polynucleotides comprising a polynucleotide sequence described by				, AL042989,
HCROW80 876942 Preferably excluded from the polynucleotides comprising a nucleotide sequence described by				, AL038509,
AL037526, AL036767, AL019464, AL038520, AL036733, AL037027, AR066494, AR060234, AR066492 Preferably excluded from the AB026436, AR054110, present invention are one or more polynucleotides comprising a nucleotide sequence described by				_
AL119464, AL038520, AL036733, AL037027, AR066494, AR060234, AR066494, AR060234, AR066494 AR066494, AR060234, AR066494 AR060234, AR066494 AR060234, AR066494 AR060110, present invention are one or more polynucleotides comprising a nucleotide sequence described by				, AL036767,
HCROW80 876942 Preferably excluded from the polynucleotides comprising a nucleotide sequence described by				
HCROW80 876942 Preferably excluded from the AA330056, AA236014, present invention are one or more polynucleotides comprising a nucleotide sequence described by				_
HCROW80 876942 Preferably excluded from the AA330056, AA236014, Z98049, present invention are one or more polynucleotides comprising a nucleotide sequence described by				_
HCROW80 876942 Preferably excluded from the AA330056, AA236014, Z present invention are one or more polynucleotides comprising a nucleotide sequence described by				, AR054110,
a Jed	-	-	excluded from	, AA236014, Z
a Sed			ĭ	
sequence described	-		ď	
			sequence described	

			the general formula of a-b, where a	
			ny integer between 1 to	
			b is an integer	
			15 to 793, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1655, and where b is greater	
			than or equal to a + 14.	
1656	HLQER45	876943	Ω	AI626059, AI626106, AA826765, AI040137,
-			present invention are one or more	AA643166, AA700884, AA548726, AW361733,
			polynucleotides comprising a	AI424257, AI860448, AA580441, AI985034,
			nucleotide sequence described by	AI720331, AI720332, AI459935, AW383179,
	-		the general formula of a-b, where a	AA308449, AW383230, AW383291, AI304515,
			is any integer between 1 to 1048 of	AI084026,
			SEQ ID NO:1656, b is an integer of	AA135152, AA588817, AA588576, AW383112,
			15 to 1062, where both a and b	AW383292, AI829153, AW383143, AW016001,
			correspond to the positions of	AI802779, AW361734, AA129139, AW383175,
			nucleotide residues shown in SEQ ID	AI475415, AA834407, AI247812, AI282992,
			NO:1656, and where b is greater	AW376286, AW392915, AA502781, AA053766,
	•		than or equal to a + 14.	AA973594, AW238610, AI860189, AW084925,
				AA344804, AW363161, AA129138, AW004060,
				AW363048, AA053663, AI638684, AW024090,
				AI694258, AA159581, AA345424, AW363163, T72477
				AA933684, AA553869, T72849, AA513679, AW352403
-				
	•			2, AW0848
				X91863, X91864, E02175, U62658, D16913,
		_		AF099176, AL080126, L24896, AL137292, M30514,
				AF161699, Y10823, L13297, AL110224, A07588,
				AR068751, AL117416, AR038969, I17767, X54971,
				E02914, Y10655, AF061795, AF151685, AL050092,
				A08913,
				S7771, A08912, A08910, A08911, I49625, A08909
	_			AF090943, AF026030, I03321, A03736, AR038854,
				A18777, A08907, A08908, AL137461, AF017152,

			1
			A90832, AF016271, AL137267, AL050280, AF159148,
			AF061943, AF008439, I18355, I34392, AL080162,
			AL137550, AB007812, AJ001838, AF117959, X76228,
			AF118064, AL050024, X70685, AF118090, AF141289,
	_		AL049464, AF017437, AR0549
			AL049452, X63410, S75997, S36676, U53505,
			I52013, AF120268, E15324, AL137558, L31396,
			U68387, AL137656, AF004162, U80742, L31397,
			I00734, AF113694, AL133558, AF069506, Y09972,
			E00617, E00717, E00778, X96540, I29004, X66417,
			AL050146, AJ012582, AL137521, AF114168,
			AF145233, AL049339, AL049300, AF113676,
			I96214, AF036941, AF055917, AF115392, U57715,
			4, AF158248
			AF031147, AL049465, AL137276, X97332, AL110171,
	_		A92311, AF113019, AL137283, U55017, U92068,
			AF051325, AF176651, AJ242859, X67688, AL080158,
		1	AF205861, AL110225, Y14634, AL117394, A52563,
			AF106934, AF119358, U91329, AF057300, AF057299,
			AF113690, AF100931, Y10080, AF022813, AL137298,
			X60786, Y11254, AL049314, E12580, X52128,
			U86379, AF126488, E01314, Z37987, AL117457,
			AL050116, AL133016, X99717, AF199027, AF106657,
			E01614, E13364, AJ012755, M92439, U51587,
			U01145, AF091084, AL050277, AB026995, AF118070,
			E12579, X06146, E15582, U77351, S82852,
			AL137554, AL117585, AL122098, AF000301,
			AL133062, AL080140, AA523439, AI652347
1657 HWADQ26	876944	Preferably excluded from the	H72650, AA486265, R36338
		present invention are one or more	

			polynucleotides comprising a	
			nucleotide sequence described by	
			the general formula of a-b, where a	
			is any integer between 1 to 598 of	
			SEQ ID NO:1657, b is an integer of	
			15 to 612, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1657, and where b is greater	
			than or equal to a + 14.	
1658	HLJBJ74	876945	Preferably excluded from the	AI089472, AI201678, AA121121, AI225034,
			present invention are one or more	AA040061, AA026978, AW074127, AA588232, R75602
			polynucleotides comprising a	AI
			nucleotide sequence described by	
			the general formula of a-b, where a	AI139753, T17035, W35381, AA161140, AA398755
			is any integer between 1 to 507 of	A1623471, H96500, C02374 AL080013
			SEQ ID NO:1658, b is an integer of	R75672, W32995, AI247236, R59185,
			15 to 521, where both a and b	AI080393, T32336, AL119457
			correspond to the positions of	4, AL119324, AL043152,
_			~	AI927233, AI538885, AI590686, AI679179,
			than or equal to a + 14.	_
				AA848053, AI446628, AI824748, AI360195,
				AI610362, AI679550, AL037081, AI625464,
				8, AL042866,
				ι,
				7, AI583578,
				4,
				, AI553645,
				AW075305, AW103878, AI284515, AW087199,
				, AW051088,
				AL041928, AW268122, AI571868, AI624529,
				, AI652162,
				AW151136, AW084065, AI539771, AI922561,
				AI432644, AI584140, AI686817, AI537677,

500659,	5, AI459322, AI815232,	5, AI682891, AI500523,	١,	9, AI284517, AI500706,		9, AI521560, AI500662,	2, AW172723, AI284509,	8, AI440263, AW088899,	3, AI434256, AI866469,	9, AI554344, AI888661,	3, AI888118, AI873638,	2, AI859991, AI436429,	, AI623736,	3, AI491710, AI431307,	6, AW151451, AI610557,	5, AI242736, AI376376,	9, AW151979, AI537187,	9, AI076761, AI539707,	8, AI963846, AI885949,	9, AW089557, AI559957,	, AI469775,	, AI567953,	, AI570966, AI53719	, AW103398, AI35501	9, AI610115, AW150457,	, AW129230,	9, AI872722, AI567582,	3, AI610402, AI370812,	9, AI624693, AL046052,	3, AL047422, AI440238,	0, AI539153, AW081383,	8, AI345477, AI683497,	5, AI933992, AI582461	U77594, Y11587, AB026436	2, U49434, AF058921, L10353,
AI62790	A186646	0132	AI88777	AI923989	AI49177	AI889185	AI582912	AI889168	AI63349	AI80576	AI284513	AI538342	AI889147	AI581033	AI866786	AI431316	AI887499	AI09448	AI866608	AI63341	AI521571	AI860783	AI446495	AI056694	AI364639	AI636788	AW080379	AW088903	AI96301	AI919593	AI26958	AW080298	AI58306	26,	AF115392
AI494201,	AI493559,	AI832245,	AI538850,	AI590043,	AI445237,	AI678446,	AI539800,	AL079741,	AI866573,	AI434242,	AI500714,	AI285439,	AW089275,	AI371228,	AI440252,	AI860003,	AI828574,	AI539781,	AI702065,	AIS69309,	AI285419,	AI865320,	AW183130,	AW193139,	AI886594,	AW085786,	AI300354,	AL039456,	AI910464,	AW162194,	AI567971,	AI627893,	AI500504,	AL117568,	AF090901,
											_											-	_	_			-								-
								_																	_										

	103321, AR034821, AL137268, AL137712, AL137658,	2
	I09499, AL133049, AL133067, I89947, S83440,	
	, AF107847,	
	AF199027, AL110222,	7
	, I48978, U96683, AF04	
	0, AL133081, M27260,	-
	L049452, AL122050, AL122100,	
	A21103,	35,
	34, AL080139, AL1375	54,
	56494, X62580, Z72491, AF114818,	
	912, AL137480, A08910,	
	I33392, U42031, AI	
	6, S77771, AF032666	15,
	08908, AF031147, AF	
	AL133665, S76508	
	AF017437, AL133558, E03349, AF159615, A30910,	
	122, AL117460, AL122045	38,
	F102578, AF	99,
		ω
	847, AR019470, AF094480, AF18221	
	013, AL122110, A65341, AL133080, AL1	6
-	68233, I92592, E01314, AL023657, AL13	
	, AL122123	86,
	0, AF210052, AL137574, AF090900, A4578	
	9, I22272, AB019565,	
	F090943, X79812,	
	AL050172, A27171, S79832, AL133113, X66975,	
	7435,	07,
	AF118070, AL	38,
	AF039137, AL137660, AL050155, AL137294, Z97214,	_
	4227, AL117648,	
_	26124, AJ010277,	
	1, AF179633, AF113690, X66862,	
	8, AL080154, AF111851, Z13966,	
	AF183393, A58545, AL080137, AL133010, AL137555,	S
	AF000145, AF008439, AF081195, AR011880, E07361,	61,

				AL035458, 7	AL137300,	AL035458, AL137300, I00734, A08911, I8994	911, I89944,
				U75932, AF1	100931, X6	AF100931, X66871, U92068, A77033	B, A77033,
				A77035, A76	5337, AL13	A76337, AL133645, AL117626,	526, AL137459,
				AL133624, 1	AF106697,	AL050116, EC	AL133624, AF106697, AL050116, E00617, E00717,
					030513, A1	AF030513, A12297, AF106862, I68732,	862, I68732,
				വ	8523, A089	A58523, A08916, AF002985, AF012536,	5, AF012536,
					AF215669,	X61399, ALO	
					X80340, AI	X80340, AL117416, AR059958,	59958, AL080234,
				AF061795, 7	AL117457,	AF151685, AF15824	F158248,
				AL137665, A	AF104032,	AF104032, X96540, M92439,	439, AC004686,
				AJ001838, I	L13297, E1	L13297, E15582, AL117585,	585, X54971,
				AF185576, 4	AF026816,	E02152, Y10655, Y10823	655, Y10823,
				AF118094, 7	AL137478		
1659	HE8TT24	876946	Preferably excluded from the		AI347465,	AA741252, A3	A1672808,
			present invention are one or more	AA251469, A	AI275156,	H61853, H618	H61853, H61854, AA336646,
			polynucleotides comprising a	AA676384, 1	AI909660,	AA182632, AA082822	4082822,
			nucleotide sequence described by	AA311433, A	AA125933,	AJ238376, AJ238375	J238375,
			the general formula of a-b, where a	AJ238374, A	AF161479,	AJ238379	
			is any integer between 1 to 873 of				
			SEQ ID NO:1659, b is an integer of				
			15 to 887, where both a and b				•
			correspond to the positions of				
			nucleotide residues shown in SEQ ID				
			NO:1659, and where b is greater				
			than or equal to a + 14.	·			
0991	HSS1S63	876947	Preferably excluded from the	_	AA612688,		AI827363,
			present invention are one or more		AI432650,	AI802722, AI	AI239964,
			polynucleotides comprising a	AA701945, A	AA612922,	AI361623, N	N33537, AI301851,
			nucleotide sequence described by	AW002136, A	AI802741,	AA176363, AJ	AA576449,
			the general formula of a-b, where a	AA976265, 1	AA766161,	AA918580, AA	AA653969,
			is any integer between 1 to 833 of	AA148478, 1	AA827535,	AA808278, HS	H93495, H62703,
			SEQ ID NO:1660, b is an integer of	T17099, A1972187,	972187, NE	N51008, AW195377, N35315	377, N35315,
			15 to 847, where both a and b	AA468340, 1	AW272194,	AA932140, H.	AA932140, H27698, H18938,
			correspond to the positions of	AI242349, 1	AI218074,	AI915880, AA601068,	A601068,
			nucleotide residues shown in SEQ ID	AI263921, A	AI925918,	T95492, R95	T95492, R95678, AA287244,
			NO:1660, and where b is greater	AI916550, A	AA886254,	H26101, AA6	AA641272, AI985842,

			than or equal to a + 14.	l .
				AW299786, H28434, H21901, H21407, AL247273, T72816, H59524, T74771, AA931965, H60166, AA148477, AI767616, AI935706, AI640135, T28521, H24592, AA385649, T71664, AA835555, T72815,
), H26143, R29069, 5, D14524, E04020, AB017196
1661 HZ	H2CAA03	876949	Preferably excluded from the	AI200746, AA306947, AA679811
			present invention are one or more	
		-	ides com	
			nucleotide sequence described by	
			the general formula of a-b, where a	
			is any integer between 1 to 494 of	
			SEQ ID NO:1661, b is an integer of	
			where both a	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1661, and where b is greater	
			than or equal to a + 14.	-
Ĭ	HCROI77	876952	Preferably excluded from the	AA631215, AI924992, AW079378, AA988078, AI820581
			present invention are one or more	
		-	polynucleotides comprising a	
			nucleotide sequence described by	
	•		al formula of a-b, wher	
			is any integer between 1 to 530 of	
				-
			15 to 544, where both a and b	
		•	correspond to the positions of	
			de residues s	
			NO:1662, and where b is greater	
			than or equal to a + 14.	
1663 HZ	H2CBW39	876953	Preferably excluded from the	AA315245, AB011148, A90836
			present invention are one or more	
_			polynucleotides comprising a	

			nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 430 of SEQ ID NO:1663, b is an integer of 15 to 444, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1663, and where b is greater than or equal to a + 14.			
1664	ннвнм68	876954	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 1265 of SEQ ID NO:1664, b is an integer of 15 to 1279, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1664, and where b is greater than or equal to a + 14.	AI344224, AI524277, AA569876, AA806680, AA302627, AI077935, AA304941, AA304941, AA304941, AA304941, AA304941, AA3040441, AA30644, AA192362, AA192362, AA171987, AA1928715, AA171987, AA171987, AA171987, AA171987, AA30635, AA30635, AA308035, AA30636, AA30636, AA30636, AA319396, AA319396,	AI343252, AI763340, AI971555, AW195633, AW242690, AI949067, AI949493, AI831556, AI589614, AW118064, AW294645, AW022953, AW068609, AA773062, AA461578, AI350493, AA661535, AI914032, AI350493, AA64527, AI433117, AI475606, AI375626, AI307282, AA814665, AA805929, AA622783, N40708, AI355690, N29617, AA630457, AI184753, AA251540, AI769738, AI584155, AI040830, AW392440, N6235, AA66293, N41617, AA058804, AA167231, AA167230, R66016, AI143758, AA66945, AA60290, AA251498, AI868406, R66015, AA172303, AA570042, AW40136 AW007103, AA657969, AA635112, AA330367, AW402028, AI219231, AI630129, AA130522, AA344392, T98790, N45715, AA569886, J02645,	AI763340, AI971555, AW242690, AI949067, AI831556, AI589614, AW294645, AW022953, AA773062, AA461578, AA661535, AI914032, AA661535, AI914032, AA65227, AI433117, AI375626, AI307282, AA805929, AA622783, 355690, N29617, AA630457, AA251540, AI769738, AA051540, AI769738, AI094496, AI143758, AA669452, AI094496, AI219343, AA857867, T98791, AA130523, AA657969, AA635112, AA657969, AA635112, AA130522, AA344392, 5715, AA569886, J02645,
1665	HSYBF36	876957	Preferably excluded from the present invention are one or more	AI341667, AA031711,	AA180986, AI341558, AI694268, AI469856,	AI093197, N63041, N50125,

			polynucleotides comprising a	AI478279,	AI150599, AI597740, AI985206,	
			nucleotide sequence described by	AI671591,	.741942, AA037642,	AI952374,
			the general formula of a-b, where a	AA180865,	AA031648, AI800796, AA436065,	
				AA129939,	AW002265, AI074205, AI056532,	
				AI656721,	ς,	W00519,
			15 to 2509, where both a and b	AA446926,	AA043021, AA830493, AI655558,	
			correspond to the positions of	AI769027,	AA443349, AI095056, AA917703, W93	W93307,
			nucleotide residues shown in SEQ ID	AA526333,		_
			NO:1665, and where b is greater	AA101851,	AW139517, AI128702, AI276137,	
			than or equal to a + 14.	AA873711,	N98234, W76109, AI631104, AA856832	832,
				W92810, A	AA042939, H87505, AA129938, AI688779,	779,
				AA693329,		AA037641,
				AI186390,	T74071, AA031685, AA037500, R82703,	703,
				AA037234,	AW380430, AA985191, R82654, H87506,	506,
				AA938640,	AI926907, AI916503, AI696069,	
				AW140052,	1671894,	AW057528,
				AI695458,	AA046964, AA725452, AI968837,	
				AA917824,	AA054749, F10070, AA917678, AA683581	83581,
				AA937814,	AI932475, AI984598, AA046963,	
				AA053281,	AI801723, AI499751, AA085888,	
				AA031686,	AI074981, AI279953, AI809560,	
				AF038662,	AB024436, AF022367, AF142672	
9991	HWMCE91	876958	Preferably excluded from the	AA890722,	AI695176, AI223269, W15428, AI678286	78286,
			present invention are one or more	AW449557,	AI344351, AW129566, AW083717	
			g			
-			nucleotide sequence described by			
			the general formula of a-b, where a			
			SEQ ID NO:1666, b is an integer of			
			15 to 421, where both a and b			
			correspond to the positions of			
			nucleotide residues shown in SEQ ID			
			NO:1666, and where b is greater			
			than or equal to a + 14.			
1991	HUVFJ36	876959	Preferably excluded from the	AI923735		
			present invention are one or more			

,			polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 511 of SEQ ID NO:1667, b is an integer of 15 to 525, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID				
	107.17	,	and where begual to a +	0 7 3 5 0 0 0 3 4	02.03.5MK	7 7 9 9 9 9 7 7	71630015
8991	HLYBU84	876961	Preferably excluded from the present invention are one or more	AW00/548, AW365081,	AW369/50, AI817246,	A1908457, A1686944,	A1630915, AW162565,
-			polynucleotides comprising a nucleotide sequence described by	AA534893, AA621824,	AA033782, AA176242,	AA599322, AA483552,	AI096489, AA588407,
			the general formula of a-b, where a	AI862878,	AA427425,	10	AA412220,
				AA243477,	W94878, AI	AI460031, N9	N95605, AA470032,
			SEQ ID NO:1668, b is an integer of 15 to 1349, where both a and b	AA677651, AA523380,	A1148140, AI434640,	AA902530, AW026082,	AAS//431, AIS73043,
			correspond to the positions of	AI129794,	AW009274,	AA554102,	AA700766,
			nucleotide residues shown in SEQ ID	AW292794,	AI673429,	AW160961,	AW026393,
			NO:1668, and where b is greater	AW272201,	AA156869,	AA075534,	AI802460,
			than or equal to a + 14.	AA643550,	AA075634,	AI086037,	AI434128,
				AA432191,	AI934640,	AA936148,	AA832390,
				AA043287,	AI075001,	AW009314,	AA830134,
				AA769386,	AI370761,	AA075581,	
				AW337458,	AA553892,	AW380901,	
				A1613297,	AA4311/1,	AW190498,	F367/3, AA1/6143
				AT355815.	W93408. A	AA417790. R3	R37629. AI538237.
				AA190514,			AA191034, H29313,
				AW057939,	_		AA306868,
				AI016135,	AI015828,	T15760, RC	R07498, AI587586,
				AA043626,	AI034090,	_	AA083325, AA553691
				AI383781,	F21581, AA156870		_
				AA316341,	AA417694,	W25045, Al	AI147345, AI418700,
				AI202543,	AA319535,	AA933690,	R07551, T60037,

				AA376766 D19678	C15100 20020144 30111544 8678
				10 100 0 CE	'SOBBETTO' WHISHOUS'
	_			₹	, F33909, AA243536, RO
				٦,	3, W28836,
					4083438,
				_	T23201,
				AI142352, AI	i, AI762052
				AW026079, HO	H01393, R76588, AI086242, AA77753,
				AA258556, AA	AA782087, AI651923, AI306436,
			•	AA946836, AA	AA946830, AW139820, AA946595,
				_	
				7,	AA865328, T86736, AA459999, AA701556,
				_	AI188276, AI000875, AA599243, N32426,
				AI023878, AW	AI088920,
				AA126805, AI	AI800579, U20272, D32257, U14134,
+				,	AC006045
MH 6991	HWLMK6	876963	bl	T86558, R745	R74597, AA495751, AI204352, N56848,
,	S		present invention are one or more		AA460093
			polynucleotides comprising a	AA693860, R9	R97459, AI806458, R97416, AA164861,
			nucleotide sequence described by	AI241618, AA	AA362800
			the general formula of a-b, where a	AA203546, AA	N35933,
_				AI239984, AI	
			SEQ ID NO:1669, b is an integer of		
			15 to 486, where both a and b	- >	
_		_	correspond to the positions of		
			nucleotide residues shown in SEQ ID		
		_		-	
+			than or equal to a + 14.		
1670 HW	HWLPY93	876964	ጚ .	1	AI379875, AA403186, AW069343,
			present invention are one or more		AW069233, AA534411, AA181432,
			otides comp	AA032182, AIS	AI935567, AI376398, AI089572,
			nucleotide sequence described by	AI452747, AIE	AI803472, AA447447, AA236374,
			the general formula of a-b, where a	AA128133, AA4	AA477274, AI038660, AA477275,
			is any integer between 1 to 1943 of	AI002572, AA2	AA233880, AA447446, AA181371,
			SEQ ID NO:1670, b is an integer of	AW130668, AI7	AI769036, C03202, AI277470, W07713,
			both a and	AA715421, AA1	AA126867, AI680552, AA404675,
			correspond to the positions of	AA126195, C04	C04150, F30780, AA235347 AA192944

AA421799, AA024985, N80591, D79794, F37772, AA127217, AA027110, Z36263, AI925660, F35592, AW263312, AI139845, AA247376, AI038015, AI128210, AA193137, AL119598, AA249326, AA629114, F31719, AA232826, AA729266, AI193315, AA249762, AW373642, AW373769, AI375939, AI383560, T29636, AW391401, AF114264, AF056035, AF056034, S67069	W05557. AA278474, AA485179	AI380296, AW206501, AI393559, AI369479, AI362907, AI125368, AW272471, AW136950, AW273903, U46350, U46345, AF166331, M60329, AJ272227, X86395, X86396	246094
nucleotide residues shown in SEQ ID NO:1670, and where b is greater than or equal to a + 14.	preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 801 of SEQ ID NO:1671, b is an integer of 15 to 815, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1671, and where b is greater that		120 ल ला
	876965	876966	876967
	HWMBV3	нсрме16	HCRQM25
	1671	1672	1673

	AA863064, AI637610, AA075674, AA075545, AA206591	AI032744, Z60017	AA694142, AA815120, AA749173, AI005429
the general formula of a-b, where a is any integer between 1 to 577 of SEQ ID NO:1673, b is an integer of 15 to 591, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1673, and where b is greater than or equal to a + 14.	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 602 of SEQ ID NO:1674, b is an integer of 15 to 616, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1674, and where b is greater than or equal to a + 14.	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 653 of SEQ ID NO:1675, b is an integer of 15 to 667, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1675, and where b is greater than or equal to a + 14.	
	876968	876969	876971
	HWMBV7	HCRŲK24	HWLOK80
	1674	6/01	1676

			the general formula of a-b, where a is any integer between 1 to 817 of SEQ ID NO:1676, b is an integer of 15 to 831, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1676, and where b is greater than or equal to a + 14.				
1677	HNTBD04	876975	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 1305 of SEQ ID NO:1677, b is an integer of 15 to 1319, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1677, and where b is greater than or equal to a + 14.	AI379864, AI806491, AI343481, AA281624, AA844487, AI380997, AA465447, AA465447, AM152574, AM152574,	AI081896, AI378805, AI083547, AI379105, AA422096, AA583293, AA581543, AA581543, AA581543, AA581543, AA581543, AA581543, AA581543,	AIO81896, AW131833, AW170478 AI378805, AI709093, AI491963 AI083547, AA411203, AI718197 AI379105, AI379556, AI361971 AA422096, AI493410, AW370895, AA583293, W04273, AW370895, AA281683, AA890322, AI671250 AA581543, H68367, H68369, AA I40124, R36504, T10779, R832 AI239994, AI333199, AW183647 L48692	5, AW131833, AW170478, 6, AI709093, AI491963, 7, AA411203, AI718197, 8, AI379556, AI361971, 9, AI493410, AW370896, 1, W04273, AW370895, H50534, 1, AA890322, AI671250, 1, H68367, H68369, AA338712, 1, H68367, H68369, AA338712, 1, AI333199, AW183647,
1678	НWLUV59	876976		AI631843, AW370191,	AI684260, AW291703, AJ224747,	AI351574, AW300604, AJ224748,	R98436, H51098, AW194814, AJ001306
1679	HSUSF13	876977	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by	AI085974, AI989948, AW150311, AI832505,	AI858091, AI934584, AI692995, AI922557,	AI720077, AW117525, AI815035, AW069468,	AW072390, AW237303, AW102807, AA446165,

R68830, R21973, AW366386, D AA322178, AA975143, AA09607 6, AI459355, AW367977, W3144 9, AA382270, AA459696, R5741 3, AI554821, AI686576, AI537 1, AI624548, AI868204, AI955 4, AI818353, AI089970, AI5881 0, AI569975, AI866469, AI440 4, AI621341, AI609409, AI458 9, AW008779, AI950892, AI927 4, AI538692, AI610690, AI783 1, AI866801, AW262042, AI800 8, AI538850, AL036901, AW18	R78950, H26464, AI300644, AA642011, AA508205, AA508225, AW235801, AA649284, R24391, AA508374, AA035658, AA301832, AA296525, R21974, H88611, AA506194, AA370945, T90836, AI025235, H88612, AA055963, AA857378, R67525, AA018277, AI828914, R24281, H98539, AA37106, AA374691, T85743,	the general formula of a-b, where a hw3/7667, A1342228, AW295915, AA843597, is any integer between 1 to 1112 of AA031368, AA031369, AA506182, AI338064, AA031369, b is an integer of AW002066, A1128919, A1083953, AW367975, N27866, 15 to 1126, where both a and b correspond to the positions of A1953830, AA976702, A1750786, A1366199, N0:1679, and where b is greater han or equal to a + 14. NO:1679, and where b is greater hw865102, N92750, A1142994, W46779, AA044355, than or equal to a + 14. W78040, A2017375, R68943, W46978, N20969, A1750787, AA102449, H28051, W32033, N40269, N30984, R67524, AW367978, AA883585, AA725372, H84840, AW074611, R70575, AA883585, AA725372,
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8,	259, AW410259,	191, AW198090,	0	ω,	·	_		AI3669		A18862	383,	123,			9087, AW082033,	3032, AF125535, Z92846	A83556, I48978,	13, A77035, AC007298,	AF081	I89947, AF08794	, AL137459, E06743,	AF098162, AL133665,	02	AL049283, AL050024, X65873,		Ă	'n.		AF032666, X82434, AL122049,	AF030513, X53587, AC004383,		32, AL137476, X81464,	9382, I26207, X84990,	7457, AL117435,
1, AI560023, AA6	177, AIS69309, AL13425	3, AL047100,	1, AIS67944,	3, AI474646,	07, AA809974,	7, AL036923, AA470	AI433157, AI654	AI520785,	AI355779,	8, AW087207, AW16	56, AI635492, AW10	10, AI863382, AI872	6, AW151766,	AI798456,	AI522052,	724, AI859991, AI573032,	01145, AL08	AL035458, AC005291, A77033, A77035,	4, AF081195, U95739,	AL050138,	U72620, A7	AL137480,	AL137558,	AF126247,	277, A08910, A58524, A58523,	51970, AI		F151685, AF	AL133075,	AL133568,		093, Z97214, AF104032,	AL133080,	100, AL137529, AL117457,
AI62394	AW07117	AI70207	AW14931	AI61291	AA7153	AI860	AI81932	AI539771,	AA835	AW05108	AW1611	AI6904	AL03898	AW07399	AW022682,	AW10472	00076	AL035	M81784,	A91162,	AL050149,	AL110222,	AF100931,	AF061943,	AL050277,	I4897	AL096744,	U6280	AR038854,	AL133640,	AF097996,	AL122093,	AF078844,	AL122100,
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				AL137488, Z35309, AL133560, I89931, AB016226, AL133557, AF184965, AR034821, AF008439, AL137463, I49625, A08907, AL122110, X53777,
				, AL110280, A08916, AF125948, AF09
-				J 4.
				2, A23630, A18777, AI
	_			, ALO49347,
				A08911, AL122121, AF113691, AL137560, AL137538, AF090901, X93495, AL133031, I96214, AL080159,
				6, AF090903, AL133016, I09499, AL1220
				AF061981, AL080148, S76508, AL122123, AL050366,
				0, AL137627, AF113019, AL1335
				0, I33392, AF113699, Y13350, AL13308
	-			3, AF111849,
	_			AL110225, S68736, AL122118
	-			A18788, I89934, AL080110, AF09108
	_			AF031903, E05822, AF111851, U35146, AF183393,
				A21103, Y10655, S75997, L13297, S36676, AL122111
0891	H2CBE41	876978	Preferably excluded from the	AI032392,
	,		present invention are one or more	AI99234
			polynucleotides comprising a	Z21538, D20524,
		_	nucleotide sequence described by	3, D59889, D80133, D80
			the general formula of a-b, where a	C14331, D80248, D81030, D59859, D
	-			, D80195, C15076, D80269,
			:1680, b is an	D59619, D80210, D51799, D80391,
			15 to 630, where both a and b	D80240, D80253, D59787, D80227,
			ס	D57483,
		- 	nucleotide residues shown in SEQ ID	D80038, AA305409, D80193, D59610,
			NO:1680, and where b is greater	C14389, D51060, D80378, C14429, D80024, D80366,
			than or equal to a + 14.	AA305578, D51022, D59373, D80045, C75259,
				3, D80268, T03269, C14014, D596
				186, AW375405, AW360844, D80014, D8013
				AW179328, AW177501, AW177511, D51213, D80247,

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D80302		D80439, AW177505	D81111,	AW352117,	D59503,	AW179019		AW177456		D51103, AW352174	7,			05856,	W177728	57774,	010,	582,	4,	4,	C14973,	51097,	957,	60214,	452,	W378542	094,	, 969	67155,	78	3,	R02520	8,	52	190.
AW360817,	AW179332,	439, A	,6, D81				, 79673,		177733,	1103, A	.4407, D51759, D80157, AW17901	178914,	AW378525, AW352163, AI910186,	D80168, AI905856,	AW177722, AW17772	2	92, D6001	3, Z2158	D80949, C1434	D58246, D59474	16, C14	AA514184, D5109	C16955, C14957,	AI525912, D6021	T03048, Z33452,	F13796, AW378	AF154840, AF125393, U57094,	A62298, AJ132110, AF058696,	1614, X	A25909, Y12724, A67220, D897	AR00844	5808, A	4R06648		92, A43
			AW377676,	, D5810	4W17890	, AW178	51, AW3	, AW176	34, AWI	71, D51	0157, 7	12, AW1	63, AIS	, D8016	, AW177	C14298, AIS	, C0305	AI525923,	D80949,	D58246,					, T0304	15, F1	AF1253	132110,	59, D34	2724, 1	D88547, 1	, AR016	0133, 1		, A43192
AW352170,	AW352171,	AW17890	C05695, 1	AW360841, D58101,	D51250, AW178909,	AW178907, AW178754,	AW3696	179020	AW3608	AW178971,	59, D8	AW179012, AW17891	AW3521	378539	178911	8540,	AW352120, C03092,	H67854,			AA285331,	D51079,	D51221	D59551,	H67858,	AI525215,	4840,	98, AJ	AB0288	17 , eo		X82626	.28, IS	A45456,	X09669
AW366296,	AW378534,	AW179023, AW178905,						F13647, AW179020, AW176467,	AW178980, AW360834, AW17773	AW178908,	7, D517	AW179009,	78525,	117, AW	774, AW	T48593, AW378540,	781, AW	2,	0			533,	AW178986,	AI535686,	AW179013,			5, A622	18138,	2, A259	5, A94995,	02449,	2, I501	AR060138,	AR038669,
١.			5, D800	2, C142	AW1789	AW177731,	_			8	C14407	4, AW17	3, AW3	1, T114	AW178			AA80912	AW36795	7, D59	AI52522	4, AW378					U38654,	A84916,	8, ARO	D2602	A82595,	5, AB0	15013		
AW378532	AW375406	AW377672,	AW178775,	AW178762, C14227,	D80134, AW178906,	D58253,	AW179018	AW352158,	AW179329,	AW37852	T02974, C1	AW179004,	AW378543,	AI557751, T11417, AW378539,	T03116, AW178774, AW178911,	D59653,	D45260,	H67866,	D52291,	AI525917, D59317,	D80258,	AW177734,	AW167716,	AI525920,	AI525235,	AI525242,	C05763,	A62300,	AR008278, AR018138, AB028859, D34614, X6715	Y17188, D26022,	A78862,	AR060385, AB002449, X82626, AR016808, AR025207,	I50126, I50132, I50128, I50133, AR066488	AR016514,	AR054175
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90, AR, AB, X6, X6, D8, 11		AA426499, AW081325, AI985955, AW021040, AI160194, N51691, AI139313, AI378674, AA622963, AI160194, N51691, AI139313, AI378674, AA622963, AI624270, AI656023, AI418379, AI091497, AA009944, AA418983, AI336531, AI394274, AA857944, C15793, AI214264, AI277517, AI346314, N47105, AI361996, C16060, AW192963, D57940, AI536992, AI304548, AA918156, C16528, N40979, N67845, AA393695, AA857656, AI659750, H95189, AI493625, C16468, D56642, AI094425, AA552961, AI080394, R81446, AW439682, N51633, D56627, D56835, N44986, H88689, AI589928, AA379627, D56835, N44986, H88648, C16043, D57541, D57973, AA328571, D57430, AA360724, AI089758, C16179, C16087, D79736, AI445344, D56588, AI218414, R69853, AA056022, AI333062, AI004951,
	preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 598 of SEQ ID NO:1681, b is an integer of 15 to 612, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1681, and where b is greater than or equal to a + 14.	O A A O A Z A O A .
	876980	876981
	HWLFY03	HE2JX48
	1891	1682

AIG10402, AIZ81837, AN17063 D45889, A74912, I89947, I48908913, I89931, D13542, I4897 AF091084, L31396, AL133640, AF104032, AL122049, AF11301 AF113677, AL050116, AF10686 AL122098, A08910, AL050277, AL113699, AF113699, AF113699, AF113699, AF113699, AF13694, AF12594 AF177401, U35846, AL049283, AF177401, U35846, AL049283, AF12594 AF137271, AL049382, AL05014 AL37271, AL049382, AL05014 AL37271, AL049382, AL05014 AF136990, AF113699, AF113699, AF113676, AF113694, A58523, AF113676, AF118064, A58523, AF113676, AF118064, A58523, AF113676, AF118064, AF11851, AL133560, AF11880, E07108, AL110280, AL080137, I03321, AC049464, U91329, E15569, AC0763, A12297, X93495, AL13771, AL037521, AL0801275, AL137521, AL080127, X60012755, AL133067, AF133067, AC0816, AJ012755, AL133067, AL131314, AL133067, AL133067, AL133067, AL133067, AL133067, AL133067,	TOLISTIC CTTIBCIE TAFOABIT BACOATT
D45889, A74912, 1899 08913, 189931, D13542 AF091084, L31396, AL AF104032, AL122049, AF113677, AL050116, AL122098, A08910, AL AF113699, AL137459, AF113691, AL080060, L110196, AL137463, AL AL050149, AL080124, AL137271, AL049938, AF177401, U35846, AL AF177401, U35846, AL AF177401, U35846, AL AF177401, U35846, AL AF177401, AS84990, AP AF113676, AF1118064, AR059958, S68736, AF AF113676, AF118064, AR133072, X82434, AF AL133072, X82434, AF AL133072, AR011880, AF118094, AL080137, AF18094, AL080137, AF18094, AL080137, AF18094, AL080137, AF18094, AL0801329, E1 A93016, X63574, U679 AL110280, AL110225, L137521, AL049300, Y1 196540, AF119337, AL08	AI610402, AI281837,
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				Z72491, M30514, AF177767, AL122118, X53587,
		_		AL080074, AL137300, AL137533, AF106827,
		_		AC002464, AF106657, AF008439, AR020905,
				AR013797, A90832, L30117, I17767, E08631,
				AF095901, E04233, U68387, I09499, AF139986,
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				AL122100
1683	HNFHD27	876983	Preferably excluded from the	AI742835, AI469703, R98751, R83167, AI538038,
			present invention are one or more	AI215412, T96765, AA206614, R93713, AI678748
			polynucleotides comprising a	
			nucleotide sequence described by	
			the general formula of a-b, where a	
			is any integer between 1 to 1000 of	
			SEQ ID NO:1683, b is an integer of	
			15 to 1014, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1683, and where b is greater	
			than or equal to a + 14.	
1684	HWLXS11	876984	Preferably excluded from the	AI692881, AI240606
			present invention are one or more	
			polynucleotides comprising a	
			nucleotide sequence described by	
			the general formula of a-b, where a	
			is any integer between 1 to 417 of	

			O:1684, b is an 1, where both a nd to the positide residues show				
			NO:1684, and where b is greater than or equal to a + 14.				
1685	HCRPG94	876985	1.0.	AA307658,	AW381667,	AW295050,	AI525535,
				AF095791,	AF220152		
			polynucleotides comprising a				
			nucleotide sequence described by			•	
			, where both a and b				
			correspond to the positions of				
			nucleotide residues shown in SEQ ID	_			
. —			NO:1685, and where b is greater				
			equal to a + 14.				
9891	HCUG073	876987	Preferably excluded from the	AI581133,	AI183335,	AI591306,	AI859797,
			present invention are one or more	AI474090,	AA757640,	AI076898,	AI559591,
			polynucleotides comprising a	AA457735,	AW173564,		AA480846,
			nucleotide sequence described by	AA767766,	AI526090,	AI392866,	AA723065,
			the general formula of a-b, where a	AA939140,	R52542, A	R52542, AW103638, AA766199,	4766199, AA757573,
			is any integer between 1 to 908 of	AI591339,	AI910407,		W47118, AW020710,
			SEQ ID NO:1686, b is an integer of	AA580663,	AL039858,	AA708505,	AI002285,
			15 to 922, where both a and b	AW090087,	AA641818,	N63128, AI440263,	[440263, AL040827,
			correspond to the positions of	AI889256,	AA939199,	AI866465,	AI401697,
-			nucleotide residues shown in SEQ ID	AW263804,	AI538850,	AI688848,	AL120853,
			NO:1686, and where b is greater	AI886440,	AI859782,	AW161156,	AA557132,
			than or equal to a + 14.	AIS67961,	AI801325,	AW020373,	AI587000,
				AW020397,	AI624950,	AIS00714,	AA056265,
				AW020693,	AI581033,	AI961414,	T99953, AI918554,
				AW167918,	N99092, A	AI619513, A.	AI345005, AL041016,
				AI340627,	AI570861,	AI889147,	AI582932,
	-			AL121564,	AI685798,	AI698391,	AI345014,
				AI538564,	AI915291,	AW152182,	AA420722,

_												 		_													 							
	AW161579, AI471909, AI923989, AI284517,	ς,	AI811192, AI917994, AI473536, AI340982,	AW079432, AA857847, AL049048, AI866469,	AW151979, AA741027, AI371251, AI859991,	AI884318, AI440238, AI624245, AI568061,	AW075382, AI923750, AI348854, W74529, AI866573,	AA042949, AA5027	AW191003, AW071380, AL036923, AI334893, J05272,	AC007283, U00978, A91160, A91162, I48978,	Y10080, X06146, A21101, I52013, AF125948,	A08909,	A58523,	I30339,	4	AF087943, A07647, U42766, AF124435, AL122045,	, AF1130	78, A	1, A08907	E04233, A08913, AL137459, AF146568, U72621,	, A18777, I89931,	D16301,	AL050149,	AJ000937, AL137640, AL049430,	I46765, AL122100, AJ003118, AL117587, AL050280,	AF026124, AF106945	AF100931, Y16645, S36676,	A77035, AL080159, AF143957,	Z37987, AL117457, Y14314, AL080156, AR038969,	8, AF090901, AL0801	L04504, AJ012755, I89947, A17115, A18079,	A15345, AL080124, X62580, AL049382, X63162,	58, AF090903, AL0501	AF061981, I32738, AB030279, AL080163, AL133112,
												 																				-		
											•			_											-		 							

				AL137267,	I68732, D8	D83032, L13297,	97, A08916,
				AF031903,	0	, AL133568, 1	AL110225
				AL122123,	M80340, AC	AC004200, AF	179633,
				X81464, AI	AL137627, AR	AR013797, AF	AF207750, AF113690,
				AF017437,	X66871, AL		AL049283, I33392,
				AF051325,	AL049464,	L30117, M8	Ω2
				AF199027,	S	_	AL133569, A52563,
				AL137527,	Y07905, AF	AF139986, AR	
				AL137665,	AF061943,	U72620,	AL137550, AL137539,
				AL117648,	AL049347,	AF0388	47, Y10936, A90844,
	•			AL137560,	E02349, AL110296,	110296, AF	AF090886, AL096744,
				I25049, I	I25048, AF177401, X86693, AF03	7401, X866	93, AF039138,
				AF039137,	AL117394,	AL133010, AF112208	AF112208,
				AJ005690,		X72889, AS	AL137479, X72889, A90832, AL133665,
				I80062, E	E02152, I795	I79595, AF002985,	185, \$75997,
				AF113694,	X82434, AF	AF119336, AF	AF090943, AB031064,
				AF069506,	AL133624,	AL133624, AL110221,	X54971, U57352,
				AF016271,	AL117443,	AL117443, AL137641,	AL137480,
		_		AL049452,	I29004, X66417,	117, AL11	AL110159, AL133560,
				S61953, Z	Z48796, AF028823	_	AL137283, I28326,
				AF067728,	X87582, U6	U67958, A93350,	350, AL137529,
				E07108			
1687	HPMDD49	876989	Preferably excluded from the	AL134806,	AW408278,	AW382759,	AA315582, N43819,
·				AW393044,	AA310712,	AA321625,	N26436, AW393061,
			polynucleotides comprising a	AA089543,	AA740922,	AW364275,	AW402662,
			nucleotide sequence described by	AA281391,	AI540961,	AI271339,	D25278
			the general formula of a-b, where a				
			is any integer between 1 to 1582 of				
			SEQ ID NO:1687, b is an integer of				
-			15 to 1596, where both a and b				
•			correspond to the positions of			Ť	
			nucleotide residues shown in SEQ ID				
			NO:1687, and where b is greater				
			equal to a + 14.				
1688	HCNSF23	876990	bly excluded from the	AI394043,	AI198754,	AI198189,	AA969930,
			present invention are one or more	AI739036,	AI268413,	AA861762,	AI222281,

	1		polynucleotides comprising a	AA883969,	AI312584,	AW197737,	AI337319,	W60319,
			nucleotide sequence described by	A1476496,				AW418714
			the general formula of a-b, where a					
			teger between 1 to 315	· ·				
			SEQ ID NO:1688, b is an integer of					-
			15 to 329, where both a and b					
			correspond to the positions of					
			nucleotide residues shown in SEQ ID				•	
			NO:1688, and where b is greater					
			than or equal to a + 14.					
1689	HKDBC15	876991	Preferably excluded from the	AI862551,	AI765006,	AI917375,	AI972770,	
			present invention are one or more	AA552639,	AI218562,	AI768706,	W65408, A	AI350781,
			polynucleotides comprising a	AI640306,	AA574291,	AA468717,	AI307307,	
			nucleotide sequence described by	AA055447,	AA514669,	AA574359,	AA516276,	
			the general formula of a-b, where a	AI658818,	AI886513,	AW104092,	AI056398,	
			is any integer between 1 to 1259 of	AW291148,	AW026517,	AI537287,	AI493566,	
			SEQ ID NO:1689, b is an integer of	AI420453,	AI962537,	AA468798,	AA477076,	
			15 to 1273, where both a and b	AA055446,	W61322, A	AI669652		
			correspond to the positions of					
			nucleotide residues shown in SEQ ID					
			NO:1689, and where b is greater					
1690	HSIGM23	876992	Preferably excluded from the	AA504588,	AL138384,	R78587, R		AA236105,
			present invention are one or more	AI367325,	R26008, H	R26008, H25950, AI359774,		AI222758,
			polynucleotides comprising a	AI285942,	AI499688,	AW072370,	AW072370, AI042411,	
			nucleotide sequence described by	AA928406,	AI817207,		AW016387,	
		. =	the general formula of a-b, where a	AI082279,	AI073537,		R78588, R63806, AA40554	05549
		_	is any integer between 1 to 1006 of					
			SEQ ID NO:1690, b is an integer of					
			15 to 1020, where both a and b					•
			correspond to the positions of					
			nucleotide residues shown in SEQ ID					
			NO:1690, and where b is greater					
			than or equal to a + 14.					
1691	HCQBN43	876993	Preferably excluded from the	AI688703,	AI761358,	AI813766,	AW182487,	
	,		present invention are one or more	AI829360,	AI380125,	AI890417,	AW377304,	

			פ אמים יואמרים מילי דוסי רביייין	AT934593	AW377372.	AW377334.	AW377268,	
			polymertectures comprising a	AW375342,	AW377315,	AI357827,	AW377285,	
			the general formula of a-b, where a	AW377266,	AA305061,	AI559533,	AW377387,	
			general remarks of a figure of the second of	AW377252,	AW377383,	AW377255,	AI283201,	
			SEO ID NO: 1691, b is an integer of	AI286089,	AW377339,	AW377240,	AW377223,	
			15 to 1636, where both a and b	AA515982,	AI343596,	AI475146,	AW193361,	
			correspond to the positions of	AW377246,	AA579699,	AI289618,		
			nucleotide residues shown in SEO ID	AA503064,	AW377220, AI803822	AI803822,		AW375369,
			NO:1691, and where b is greater	AW351685,	T29359, A	T29359, AW377256, AW375332,	W375332, N48341	341,
			than or equal to a + 14.	AC000061,	AR016032,	111500, I	AR016032, I11500, I66544, M55131	١,
				M76128, A	83151, U20	118, A4904	M76128, A83151, U20418, A49045, AF162427,	
				166545, A	AF016950, A	AF162400, A	AF013753	
1602	HCOBO03	876994	Preferably excluded from the	AW369811,	AW014155,	AI334392,	AA664276,	
1)))		present invention are one or more	AA608594,	AA984631,	AI954111,		
			polynucleotides comprising a	AA586953,	AW194426,	AI445882,		R11024,
			nucleotide sequence described by	AA911063,	AI335787,	AI623204,	AA419568, R1	R11072,
			ral formula of a-b,	AA864381				
			is any integer between 1 to 821 of					
			SEQ ID NO:1692, b is an integer of					
			15 to 835, where both a and b					
			correspond to the positions of					
			nucleotide residues shown in SEQ ID					
			NO:1692, and where b is greater					
			than or equal to a + 14.					
1693	HCOCF85	876997	ıı					
	,		present invention are one or more					
			polynucleotides comprising a					
			nucleotide sequence described by					
			ral formula of a-b,					
			is any integer between 1 to 593 of					
			SEQ ID NO:1693, b is an integer of					
			15 to 607, where both a and b					
			correspond to the positions of					
			nucleotide residues shown in SEQ ID					
			NO:1693, and where b is greater					
			than or equal to a + 14.					

1694	HUVFS16	876998	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 1259 of SEQ ID NO:1694, b is an integer of 15 to 1273, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1694, and where b is greater than or equal to a + 14.	AA443167, AL046148, AA243821 AA243686, AA405113, AI351901 AA011361, AL043877, AB020669 AF068920, AF068921	1, AA492497, 1, AA463466, 9, AF054828,
1695	нсоврзі	877000	preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 786 of SEQ ID NO:1695, b is an integer of 15 to 800, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1695, and where b is greater than or equal to a + 14.	A1635096, AA165632, AA523697, AW166525, AA769127, AW129960, A1686907, A1768699, AW136550, A1915606, AW188763, H79957, A A1769970, AA719353, AW151462, AW418915, AA829144, AA165668, AW182418, AW102605, AA757716, C16515, AA907061, AA860897, A A1217382, A1239881, AA703100, AA577904, A1217382, A1239881, AA703100, AA577904, A1637789, N87490, N42130, A1764980, A1941067, AA649747, AA642829, R69594, A A5992380, AC006047, AP000509, AC004185, AL080317, AC005406, Z97876, AC004542, A A292380, AC003680, AC003691, AC003691, AC003691, AC003691, AC003691, AC003691, AC003691, AC003691, AC005574, AC003082, AC006023, AC002536, AC00689, AC00689, AC00689, AC00689, AC00683151, Z98257, AC0060317, AC005302, AC006832, AC006335, AC006335, AC006335, AC006335, AC006335, AC006335, AC006331, AC006332, AC006331, AC00	AA165632, AA523697, AW166525, AW129960, AI686907, AI768699, AI915606, AW188763, H79957, AI540313, AA719353, AW151462, AW418915, AA719353, AW151462, AW418915, AA719353, AW151462, AW418915, C16515, AA907061, AA860897, AI217462, AI239881, AA703100, AA577904, R21911, N87490, N42130, AI764980, AI936236, AA649747, AA642829, R69594, AA528274, AC006047, AP000509, AC004185, D84394, AC005406, Z97876, AC001604, AL030998, O04707, AC004617, AC004691, AC007319, O05908, AC003983, AL023280, AL031073, O10209, AF026254, AF026248, AF026249, AC003689, AC002094, U77841, AC004772, AC003082, AL049697, AR036572, U91328, AC002059, AJ239329, AP000688, Z98257, AC006017, AC00532, AC003087, AC007317, AC025517, Z97198, AC00385
9691	HCRMU18	877001	Preferably excluded from the present invention are one or more	AA486568, AI733856, AA077667 AA831426, AI336771, AA493546	7, AI090377, 6, AA670392,

			nolymicleotides comprising a	AT816058.	AC005914.	AL035681.	AL050307.
			de sequence describ	AC009516,	Z83826, AC	3005015, AC	Z83826, AC005015, AC007041, AC004706,
			the general formula of a-b, where a	AC005484,	AC004819,	AC007536,	AL121825,
			is any integer between 1 to 504 of	AF067844,	AP000512,	AC004962,	AC007685,
_			SEQ ID NO:1696, b is an integer of	AF109907,	AC005412,		AC005274,
			15 to 518, where both a and b	AF027390,	AC002477,	AC006487,	AC006011,
			correspond to the positions of	AL022318,	U62293, AC005730,		AC005069, U22376,
			nucleotide residues shown in SEQ ID	AC005800,	AL139054,	AL139054, AC007216, AC004150	AC004150,
			NO:1696, and where b is greater	AC000353,	Z95114, AC005754,		AL049569, AL049766,
			equal to a + 14.	AC005013,	AC005081,	AB023049,	AC006581,
				AP000558,	AP000045,	AL080243,	AC009248,
				AC005071,	AC004686,	AL109628,	AC007073,
				AC005971,	AL035461,	AL022721,	AC005164,
				AL096791,	AC005057,	D84394, Al	D84394, AL121658, AC006251,
				AC009721,	AC003663,	AC007371, AL049869	AL049869,
				AL031432,	L44140, Z	98950, ACO	Z98950, AC005520, AP000031,
				Z98946, AL022238,		AC006511, AF	AP000557, AC004668,
				AL031666,	AF207550,	AC005488,	AC005358,
				AL117694,	AC019014,	AL121603,	AL021940,
				AC007226,	AC005632,	AC005670,	AC005529,
				AC006006,	AC008115,	AC002300,	AL035086,
				AC005200,	AC004491,	AL023807,	AF200465,
				AP000116,	AC007676,	AC004149,	AF129756,
				AC007899,	AC005740,	AC006961,	AC004913, AC005088
1697	HONAN63	877002	Preferably excluded from the	AA305628,	AA308609,	AA300521,	AA356487,
			present invention are one or more	AA363124,	AB020712		
			polynucleotides comprising a				
			nucleotide sequence described by				
			the general formula of a-b, where a				
			is any integer between 1 to 530 of				
			SEQ ID NO:1697, b is an integer of				
			15 to 544, where both a and b				
_			correspond to the positions of				
			ໝ				
			NO:1697, and where b is greater				
			than or equal to a + 14.				

H O	AA987568, AL035420	AB028946
Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 518 of SEQ ID NO:1698, b is an integer of 15 to 532, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1698, and where b is greater than or equal to a + 14.	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 175 of SEQ ID NO:1699, b is an integer of 15 to 189, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1699, and where b is greater than or equal to a + 14.	preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 624 of SEQ ID NO:1700, b is an integer of 15 to 638, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1700, and where b is greater than or equal to a + 14.
877004	877005	877006
нсосибя	HCRNO79	HCRMO22
8691	6691	1700

AI400725, AI432790, AI863014, AI932794,
AL031228,
 AL137556,
AL080074, AL122098, AL137558, AF012536, I48978,
189947,
9625, E03349, A08
S77771, AR038854, AL133
06779
AF119337, AF000145, I26207, D83989,
Y08769, AL122045, I66342, AF106657, AL133010,
v
5, AF210052
 I89944, AF113689, E02253, AR059958, U96683,
 AL137640, S68736
A77033, A77035, S76508, AR000
D89079, U39656, I42402, E15569, AF032666,
AL137463, AL080060, AL137429, AL133067,
 AF132676, AF061836,
 AF090886, AL137712, AL137527, E02221, AF111112,
7215669, AI
AL133665, AF125949, A45787, AL133077, AL137658,
AL137294, AF113691
'078844, AF118070,
AL080140, S79832, AF022363, AL122121, U72620,
5341, J05032, AL13
, AF104032, I48979, AF003737, X72387,
 , AL050277
 AL133558, Y11587, U00763, X62580, AL049382,
5, AF090901
 I41145, S61953, A211
 AL080086, AF113019, AL049460, E1558
 28823, AF100931, AL122049, L19437,
AF118064, AL137478, AL122050, AL080159,

				AL133640. AL133098, X52128, AF159615, I17544,
				3, X92070,
				133075, AF061795, AF151685,
				AL096744, AJ003118, AF158248, U49434, AF061981,
				AL133568, AF146568, AL080148, AL133113,
				E13364, AF106862,
				AR019470, I33392, Z82022, AF176651, AF183393,
				AF153205, AF106697, A52563, AF139986, A08915,
				AF057300, AF057299, AL137283, AL117585, Y10080,
				AR068751, S75997, AR029490, Z72491, AL133081,
				AL049452, AL117460, L31396, I80064, AL137521,
				L31397, S78214, M92439, A15345, AL049464,
				AL117648, AF090934, AF118094, AL137557, U95114,
				AL110196, AL049466, AF118090, AL049314,
				AL080154, I03321, U58996, E06743, A90832
1703	HHPEK59	877009	Preferably excluded from the	AA149062, W55857, AI654104, N91520, AA398769,
			present invention are one or more	AL041623, AA149063, AA307763, AW450873,
			polynucleotides comprising a	AI082461, AA709060, W06955, AI079909, AI920841,
			nucleotide sequence described by	AA292830, AI268616, AA191706, AA010085, R07052,
			. formula of a-b, where	Z44437, T87013, T12757, Z40368, AA844584,
			is any integer between 1 to 1606 of	AI955471, W55858, AW135814, T52489, N48933,
			SEQ ID NO:1703, b is an integer of	T56321, N46430, AA864954, AI274165, AF027218,
			15 to 1620, where both a and b	AF027219, AF155101
			sitions of	
			nucleotide residues shown in SEQ ID	
			NO:1703, and where b is greater	
			than or equal to a + 14.	
1704	HKCTB07	877010	Preferably excluded from the	AF105020
			present invention are one or more	
			leot	
			nucleotide sequence described by	
			the general formula of a-b, where a	
			is any integer between 1 to 391 of	
			SEQ ID NO:1704, b is an integer of	
			15 to 405, where both a and b	

			correspond to the positions of					
			le residues shown in					
			NO:1704, and where b is greater					
			than or equal to a + 14.					
1705	HFPIZ22	877011	Z	AI458123,	AA770557,	AW299665,	AW236534,	
		-	present invention are one or more	AI952929,	AI340145,	AI339835,	AI650682,	
			polynucleotides comprising a	AI472033,	AA256229,	AI268229,	AA678840,	
			nucleotide sequence described by	AW190757,	AI075831,	AI631649,	AL138340,	
			the general formula of a-b, where a	AW080424,	AA293773,	AI373728,	AA704702,	
				AA677322,	AI033016,	AW204318,	AA848089,	
			SEQ ID NO:1705, b is an integer of	AI891160,	AA399568,	AA227660,	AI001981, N2	24286,
			15 to 1592, where both a and b	AA747722,	AI537348,	AW025794,	AA218733,	
			d to the po	AI865908,	H98718, H	H64686, R38180,	180, R17022,	
			de residues s	N70123, AI493281,	I493281, A	AW007482, H70397,	', AW13	908
			NO:1705, and where b is greater	AA334373, W04161,	W04161, R	R09968, AA394090,	94090, R16715	5,
			than or equal to a + 14.	T77116, W	W01375, AI690748,	90748, AWI		
					AI245731,	AI273189,	AI627988,	
				AI698391,	AI368579,	AI969655,	AW149925,	_
				AL046835,	AI690687,	AI524654,	AI289310,	
				AI868204,	AW051088,	AI869377,	AI678446,	
				AI613038,	AI590043,	AI469587,	AA464646,	
				AI589428,	AI590830,	AI863382,	AI677797,	
				AI621341,	AW149076,	AI536574,	AI538850,	
				AI921254,	AI927233,	AI568592,	AIS90423,	
				AW020397,	AI583982,	AI950892,	AL045266,	
				AI335208,	AI491775,	AI865906,	AI612913,	
				AI888208,	AI670009,	AI433157,	AI702073,	_
				AI890507,	AI682968,	AI401697,	AI538564,	
				AI445611,	AI679266,	AI913312,	AI686576,	
				AL037454,	AI627893,	AI586931,	AI872545,	
				AL037582,	AL037602,	AI815232,	AI281757,	
				AA766116,	AI537677,	AI434731,	AI635634,	•
	-			AI648454,	AI634467,	AL036802,	AI540674,	
				AL039086,	AL036673,	AI471282,	AW162194,	
_	•			AI582932,	AW148423,	AI923989,	AI583578,	
				AI866770,	AL120300,	AI890907,	AI370623,	

	AISTOLOGY, AIGHORY, AISTOLOGY, AIS
A67588, A7	A77033, A77035, AL050108, AL050138,
AF199027, A	, AL117435, A08916, A08910, AL035407,

	1 37967	פעומפטזע בעכטטטטע פנאסעסדע כככעד
	•	, ALOGAS, AFOOCEST, ALCOCTE
	AL049452,	ο <
	ALIZZOSO,	, ACUU/II4, AFUB//28, T66342 297214 AF11
	574	AF104032, AF0910
	AL049300,	L137478,
	AL133067,	, AL133640, AL137459, AF090903,
	AF177401,	U67958, AL080159,
	AL133113,	AL049283,
	AL110225,	21, X96540,
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	A58524, p	A58523, E02349, Y09972, A08912,
	AF090896,	, AL137294, AL050393, A18777, I89931,
	Y11254, A	Y11254, AJ000937, AL110221, AL117457, AL050116,
	AL049339,	, AF158248, AF090901, A03736, AF115410,
	AR011880,	, AL133637, X79812, AL050024, Z13966,
	AF061795,	, AF151685, AL137533, AL137550,
	AF061573,	, AL137292, S76508, S61953, I49625,
	AF113690,	_
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	AF118070,	, Z82022, AF100931,
	AL137557,	, AJ238278, E07108, AF
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	AR020905,	, AF113694, AF113677, S63521, AF118064,
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	AL117394,	, AL133619, A211
_	836676, 4	A93350, Y14314, AF057300, AF057299,
	AC006112,	AF090934, Y11587, AF
	X84990, p	F081197, AF081195, AF118094,
	AL050155,	U35846, AL137479, AL080124,
	AB019565,	S75997, AF113019, AL110196,
	X70685, A	NF115392, AF125948, AF125949,
	AL080140,	031, U78525
	106	, X98834, D83032, AF126247, AF082526,
	A76335	

17051	UCSEDSO	877013	Drefershly excluded from the	AT797081 AT669186 AT922708 AI400881.
2	707 10711		present invention are one or more	, AA062971, AW027338,
			ĕ	AI091639, AI627975, AI358574, AI202381,
			nucleotide sequence described by	AA255522, AW086138, AA890259, AA806628,
			the general formula of a-b, where a	AA255565, AI367251, AA088310, AA765366, D63210,
			is any integer between 1 to 1428 of	AI796381, H48099, H48098, AA720634, AL079437,
			SEQ ID NO:1706, b is an integer of	AI758780, AI911927, AW022560, AA256707,
			15 to 1442, where both a and b	AA737329, AA255588, AA877667, AA455364, AA813874
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1706, and where b is greater	
			Ũ	
1707	HCRND67	877013	Preferably excluded from the	
			present invention are one or more	AI949312, AA252030, AA521447, AW024768,
			polynucleotides comprising a	AI039260, AI962419, AI935656, AI416968,
			nucleotide sequence described by	AI361764, AA860961, AI127900, AI936802,
			the general formula of a-b, where a	AI761487, AI580311, AI917267, AW024010,
			is any integer between 1 to 794 of	AI189597, AI864624, AA131263, AI351462,
			SEQ ID NO:1707, b is an integer of	AI422420, AA904280, AI636058, AA931114,
			15 to 808, where both a and b	AA648498, AI767707, AW262532, AA191430,
			correspond to the positions of	AI312828, AA860568, N46577, AA804488, AI680207,
			nucleotide residues shown in SEQ ID	AA628794, N45139, AI694810, AA574232, AI522273,
		_	NO:1707, and where b is greater	AI362932, N46583, AA364681, H91961, N40538,
			than or equal to a + 14.	W22178, H99173, W22807, AA829581, AL046944,
				R79750, AC005325
1708	HSPA101	877014	Preferably excluded from the	
			present invention are one or more	AI457284, N35406, W49563, AA334557, R58493,
			polynucleotides comprising a	H24416, A1678442, AI791556, AA242954, R30676,
			nucleotide sequence described by	AW022665, R47185, AL031652, L41349, L13935,
			the general formula of a-b, where a	L13936, L13937, L13938, AL117633, L15556,
			is any integer between 1 to 1041 of	L18962, AF027571, AF031370, U57836
			SEQ ID NO:1708, b is an integer of	
			15 to 1055, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1708, and where b is greater	

			than or equal to a + 14.	
1709	HOSXA83	877015	Preferably excluded from the	AA100220, AI167817, AA113216, AA324768,
				AA085997, AA149087, AI493421, AA629345,
			polynucleotides comprising a	AA625949, AA149086, AA669959, AA431870,
			nucleotide sequence described by	AI866312, Z28464, AA172371, AW173386, AI183937,
			the general formula of a-b, where a	AA431871, AA262957, AL036908, AI271960, AA085643
			SEQ ID NO:1709, b is an integer of	
			15 to 1044, where both a and b	
	_		correspond to the positions of	
			nucleotide residues shown in SEQ ID	
	_		_	
			equal to a + 14.	
1710	HAVTF85	877018	Preferably excluded from the	
			present invention are one or more	AL039480, AA442561, AA858311, AI566218,
			polynucleotides comprising a	AA846839, AI583216, AI635043, AA699924,
			nucleotide sequence described by	AI192601, W69310, AI262270, AA526986, AI304664,
			the general formula of a-b, where a	AI310345, W69206, AI147372, AA973817, AI431515,
			is any integer between 1 to 881 of	AI818856, AI033497, AA983644, AW129307,
			SEQ ID NO:1710, b is an integer of	AA701244, AA926804, AA630163, AI289870,
			15 to 895, where both a and b	AA554361,
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	AI367351,
_			NO:1710, and where b is greater	AA808417
			than or equal to a + 14.	
				AA291486, R28626, W68336, H29812, R52537,
				R42369, H02369, F02630, AI686839, AA188995,
				AW236685,
				C02595, AA653337, AA883260, R43407, T29673,
				AI471055, AA190445, AI567050, AA031670,
	_			AI246665, AI658622, R33489, AI932403, AL041862,
				AI923989,
				AW071349, AL046356, AI554245, AL119748,
				AL079977, AI815232, AL046926, AL040243,
				AI434223, AL047675, AI866573, AL042628,

ALD 33 785, ALG 359 78 ALD 42744, AW151136 ALS 4821, AIS 38716 AIS 4821, AIS 38716 AIS 54821, AIS 9183 AIS 9189, AIS 9189 AIS 9189, AIS 9189 AW129106, AIS 9189 AIS 38 342, AIS 9147 AIS 38 342, AIS 9147 AIS 38 342, AIS 9189 AIS 37273, AIS 9189 AIS 37273, AIS 9189 AIS 37273, AIS 9189 AIS 38 36, AIS 9185 AIS 37273, AIS 9189 AIS 92 512, AIS 9189 AIS 92 8401, AIS 92 82 AIS 92 8401, AIS 96 90 AIS 92 93 84131 AW190042, AIS 86 863 AIS 88 9953, AIS 48 663

	I48979, AL110225, AL122049,
	I89947, AL133072,
	AL133016, A12297, AL137271, A08916, AL122050,
	A08913, A08910, A08909, AF078844, I33392,
	AF067728, AL049283, AL133080, I89931, AL050277,
	AF017152, S68736, AF146568, AL050138, I49625,
	AL122093, AL122110, Y11587, AF113689, AL137557,
	AL133560, AB019565, U91329, Y11254, AL133640,
	AL117457, AL133077, AL080124, AL133606,
	AF113677, AL137550, AL137459, E07108, AL050108,
	AJ000937, AL049314, S78214, AL133075, AL096744,
	AF113691, AL133557,
	AF125949, AL137527, AF106862, X98834, AF113019,
	F090934, AF
	I26207, U3
	7,
	1, U72620, AL080060, X72889, AL13
	AR059958, AL137538,
	AL080137, AF104032,
	7, AL0494
	E02349, E15569, A93350
	E00617, E00717, E00778, AL050146, AL137463,
	A65341, AF087943, AL049382, I42402, Z82022,
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	A58523, AL050149, AF111112,
	L050172, AL049464, L31396,
-	61943, X93495, A08912, AL137476, AL049300,
_	106/958, AFI1933/, AL13/283, AL080159, AL110197.

ALL137533, IO9499, AF AR000496, U39656, L1 S61953, A90832, Y099 E04233, Y14314, ALL11 AF185576, ALL137523, AF057299, A45787, AL ALL133067, U58996, E0 AL080074, X53587, E0 E08631, AF061573, AF ALL17440, ALL137273, I17767, U96683, X835 U68387, AR013797, X8 ALL133049, ALL117432, L05186, E12747, ALO5 AF132676, AF061836,	from the AW135340, AI908516, AW003833, AI692953, Te one or more AI693316, AW242982, AI194008, AI672260, AI497695, AW242975, N63914, AW242988, AI341520, described by AI972371, AI373504, AA705554, AI633950, of a-b, where a AI276537, AA699365, AI989919, AW204605, H11413, ten 1 to 1600 of RO0441, RA279329, AI656862, AI961706, AA455604, is an integer of AA455968, W32633, AA528280, AI702940, H85245, bit a and b T95059, H08429, F13395, T81953, F37163, AA215977, AA301556, T95155, F11101, T77655, H11389, AA279895, AW196491, AI915713, N80005, AI584140, AI890223, AI612913, AI648509, AI539687, AI537677, AW262565, AI594011, AI539687, AI537677, AW262565, AI569616, AI801766, AI610402, AW071349, AI811344, AI520785, AI680498, AI591316, AI554818, AI688872, AA225339, AI239205, AI568670, AI801744, AI810444, AI83076, AI759867
	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 1600 of SEQ ID NO:1711, b is an integer of 15 to 1614, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1711, and where b is greater than or equal to a + 14.
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965	AI44023	AI53977	AI24925	AI70107	AI56471	AW13077	AW149311	A127304	AI95066	AI49806'	AI63341	AL04115	AW09001.	AW15057	AI590120	3636	3	AI590118	AI62028	AW08103	AI09724	AI86660	213	980	AIS9002	AI624206	AI857296	AI80115	AI34900	AI57098	AL11040	AI78425	804	AI57090	AI43303
61950	AIS67846,	AI491852,	AI364788,	AI873604,	AW170635,	1779	AI538085,	AI284484,	AI868831,	AI247193,	AW088903,	AL045500,	AI270055,	AI633125,	AI536638,	AI863014,	AI282504,	AI274013,	AI648663,	AI475451,	AI434223,	AW301409,	AI475394,	S	9	AA807352,	AA470491,	AW169671,	AI445237,	AI828731,	AI564247,	AI889376,	52467	95591	AI445025,
58	AW132056,	AL040243,	AI637584,	AI859511,	AI926790,	AL134830,	AW026882,	AI702073,	AI679990,	AIS71909,	9	AI540832,	AI587143,	AI318280,	AI439745,	AW302988,	AW051258,	AI610362,	AL046944,	AI281837,	AL043981,		[AI500523,	AI270707,	AL039276,	17	AI696612,	AW274192,	AL041573,	04332	55434	AI572787,
10	AI554427,	AL119863,	AI500077,	AI559296,	AI890833,	AL045266,	AI569583,	AI433157,	AI934036,	AI475371,	AI280747,	AI280751,	AW193000,	AI627360,	AI673785,	AI274508,	AI275175,	AI537024,	AI815855,	AI568296,	AI922901,	AI702068,	AI254042,	AI687362,	965	AI801325,	AL121270,	AI500706,	AI536685,	AW151138,	AI500662,	AI499285,	W26	AI921248,	AI648454,
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AL121463, AI884469, AI648684, AI612759,
AI560099, AI064830, AA835801, AL043975,
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AL050393, AL117460, AL1174
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AL117457,
U42766, AF113690, AL133080, AL117585, AL050149,
 AF090901, A77033, A77035, AL049452, AL122093,
AL137557, AL050116,
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AF017152, AL096744, AL133016, AF078844,
E07361,
AF125949,
, A08910, AF177401,
F183393, AF125948,
, A93016, AL
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 , AB019565,
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1, AL137538, X70685, AL13764
03736, X65873, AF061943, AF0677
, AL049283, AL080159,
AF087943, AL133568, AL133072, AF111112,

HOSBX95 877020	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 516 of SEQ ID NO:1712, b is an integer of 15 to 530, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1712, and where b is greater than or equal to a + 14.	AL122049, AL137521, I09360, AR000496, U39656, 142402, E08263, E08264, AL122111, AL133067, U67958, AL110197, E15569, A93350, AL137533, AL137523, AF057300, AF057299, AF026124, AL137560, AL137480, AR013797, Y09972, AF026816, 126207, AL050172, S61953, AL137556, AL137526, 100734, E00617, E00717, E00778, U68387, E02221, 166342, A08911, Z37987, AC006371, Y14314, AR038969, A07647, AL110280, AL137429, AL080074, Z72491, AL137292, AL137476, Y10655, AF003737, U78525, AL080148, U96683, AL133104, AF100931, E06743, AF106827, AF159615, AF185576, X87582, 117767, A45787, AF061981, AL13358, AF111849, AL137488, AF162270, E08631, AL122118, Y07905, AF005922, AL122045, AF095901, AJ006417, E04233, AL133081, I09499, AL110222, L30117 AW393918, N56766
HSIFP30 877022	Preferak present polynucl nucleoti	AI678780, T98311, R10554, AF209389

		is any integer between 1 to 714 of	
		de residues show	
		than or equal to a + 14.	
1714 HE9HL05	05 877023	Preferably excluded from the	4, AI310154, N48237,
		present invention are one or more	AA333785,
		polynucleotides comprising a	T95816, AI678780, T96750, R91078, AA344220,
		nucleotide sequence described by	R09895, T74622, T68354, N49552, AA332963,
		the general formula of a-b, where a	AI023306, T71511, T95519, R92515, T60367,
		is any integer between 1 to 1581 of	AI791396, AW172723, AI815239, AI362332,
		SEQ ID NO:1714, b is an integer of	AI249946, AA665587, AW078729, AI805769,
		15 to 1595, where both a and b	AW265004, H42825, AI669639, AI608802, AW074274,
		correspond to the positions of	AI702540, AI499104, AI758816, AW263799,
		nucleotide residues shown in SEQ ID	AI886163, AI476147, AI677797, AW026633,
		NO:1714, and where b is greater	AI816956, AI677647, AI911645, AI961622,
		than or equal to a + 14.	AI250175, AA614660, AI244380, AI446124,
-	, , , , , , , , , , , , , , , , , , , 		, AI869750, AI921609,
			AA810969,
			AI446564, AI419311,
			AI628855, AI446110, AI872810, AI471424,
			AW150505, AI570195, AW150351, AW118457,
			AI694855, AI419417, AI369029, AI474427,
			AI568870, AW079656, AA088789, AI521128,
			AW168031, AI660848, AA910956, AI701948,
			AI589433, AI805385, AI591381, AI333552,
	-		AW263697, AI679622, AI683465, AI610645,
			AI952302, AI625231, AI696626, AI890714,
			AI347569, AI671638, AI560514, AW193020,
			AF209389, J04813, M18907, X12387, M14096,
			E02555, D31921, D00408, E02532, J04449, S53047,
			, M13785, AF182273, L26985, X5491
			U59378, AF109068, Y10214, M73992, Y11995,

				AF204959, AF185589, D11131, S74699, S74700.
				L35912, I12087, AF067420, A94751, U77594,
				AL137561, AC004455, AF109906, U92068, A69673,
				A69681, U89906, AF106934, AF059612, AL133645, AR068182, AL137659, AC005284, AC007370
1715	HWLMB91	877024	Preferably excluded from the	AI188270, AI742085, AI167453, AW204725, R53616,
			present invention are one or more	R48325, AA347732, AW341017, AA579588, F35057,
			polynucleotides comprising a	AA768452
			nucleotide sequence described by	
			the general formula of a-b, where a	
			is any integer between 1 to 577 of	
			SEQ ID NO:1715, b is an integer of	
			15 to 591, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1715, and where b is greater	
			equal to a + 14.	
1716	HOVEE11	877025	Ω١	AI762892, AI760766, AI174624, AW081757,
			present invention are one or more	AI824008, W94214, AI189223, AA447177, AI927354,
			polynucleotides comprising a	
			nucleotide sequence described by	W81043, AI934550, AA605197, AW390982, AI168782,
			the general formula of a-b, where a	W81079, N56763, AW374587, W72920, AI538814,
			is any integer between 1 to 1960 of	AW079505, AW137328, AA629096, AI699821, AI767317
			SEQ ID NO:1716, b is an integer of	
			15 to 1974, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1716, and where b is greater	
			than or equal to a + 14.	
1717	HCYBN69	970718	Preferably excluded from the	AA127756, AA769607, AA305740, AW403303,
			present invention are one or more	AA361909, D81026, D81030, C14389, D80522,
			polynucleotides comprising a	C15076, D80133, D80166, D80193, D80212, D59502,
			nucleotide sequence described by	D80022,
			the general formula of a-b, where a	D59787,
			is any integer between 1 to 545 of	, D80269, D80240,
			SEQ ID NO:1717, b is an integer of	D80227, D59927, D80219, D51423, D51799, D80253,

correspond nucleotide NO:1717, an than or equ	to the positions of residues shown in SEO ID	10024, D8025
nucleotio NO:1717, than or o	residues shown in SEO ID	
NO:1717, than or 6		5, C14429,
than or e	d where b is greater D802	D80268, AA305578, D510
	equal to a + 14.	D51022, C06015, AA
		AW177440, D80014, D80439, D80302, C14014,
	AW360	186, D802
	D80132,	AW375405, T02974,
	D51213,	DS9503, AW178983,
	C14227,	C14077, AW360844, D58101, AW36
	AW375406,	AW377676, D51103, AW378534,
	AW377672,	AW177501, AW179023,
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	AW178762,	8762, D80134, D51250, D51759, AW176467,
	AW352171,	
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	AW179024,	9024, AW369651, AW367967, AW352158,
	AW177505,	7505, AW360841, AW352117, AI243347,
	(AW179020,	9020, AI239543, AW178909, AW177456,
_	AW175	
	AW178908	, AW1787
	C14407,	D59653, AW17
	AW378525	, AW352163,
	AW352	AW352174, AA805151, C14298, D45260, D80168,
	AW179009	, AI905856,
	AW177	AW177722, AW177728, CO3092, D58246, AW378540,
	AW378	AW378539, AW367950, AI557751, AI525923,
	AA805	M M
	AW177508	, AI557774, T03116, D59695,
	D8094	AI535850,
	AM177	1177723,
	AIS35	AI535686, D59474, AW179011, D59551, C14973,
	AA514	AA285331, D51221, T0
	AIS25	734, D60010, D60214, AI5
	D5105	4957, C14046, AI525
	AI525	5242, AI525235, AI525222, AI525912,

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AI345993 0293, AI524526, AL047041, AL038565 6980, AI445165, AI348897, AA427700 8716, AI32339, AI34894, AI862144 2292, AW089572, AI609594, AI862144	2284, AI818683, AI273142, AW169671 2127, AW301409, AI573032, AI682971 3419, AI921248, AI669532, AI498579 5002, AI433976, AI828731, AW166970 0190, AI432969, AW102785, AI612759 9085, AI696398, AI571909, AI677796 9470, AI909697, AL045163, AI612759 9153, AA640779, AW238730, AI439745 1712, AL121463, AA572758, AI702073 1802, AI926790, AI591316, AI952360 8501, AI812107, AI8020693, AI340603 7137, AI537677, AI922901, AI349536 8501, AI812107, AI815232, AI269696 8501, AI812107, AI800453, AI340582 0433, Z99428, AI888953, AI567128, 0781, AI567993, AI349645, AW074869 0120, AW149227, AL036274, AI345131 7534, AI309401, AW103893, AI561299 6403, AI448408, AI343112, AL1212014 4517, AW071349, AI207572, AL121270 1300, AI349598, AL03664, AW075207 6456, AI648684, AW151136, AI345730 8716, AI702406, AI174394, AL041573 2992, AW089572, AI609594, AI862144 2146, AI312339, AIS84131, AI269862 6549, AW086113, AI869367, AI520785	2284, AI818683, AI273142, AW169571 2127, AW301409, AI573032, AI682971 3419, AI921248, AI469532, AI498579 5002, AI433976, AI828731, AW166970 0190, AI432969, AW102785, AI612759 9085, AI696398, AI571909, AI677796 9470, AI909697, AL045163, AI612759 9153, AA640779, AW238730, AI439745 1712, AL121463, AA572758, AI702073 6802, AI696396, AI591316, AI952360 8220, AI654750, AW020693, AI340603 7137, AI537677, AI815232, AI269696 8501, AI812107, AI800453, AI340582 0433, Z99428, AI888953, AI340582 0433, Z99428, AI888953, AI340582 0433, Z99401, AW103893, AI361299 6403, AM149227, AL036274, AI345131 7534, AI309401, AW103893, AI561299 6403, AI448408, AI343112, AL1212014 4427, AL036396, AI53664, AW075207 6456, AI64864, AW151136, AI3457700 8716, AI702406, AI174394, AL041573 2320, AL038605, AI610690, AI500077 2392, AW089572, AI609594, AI862144 2146, AI312339, AIS6311, AN26855 6549, AW086113, AI869367, AI520785 7396, AI610307, AW268251, AW268253

AI636585, AI439762, AL036631, AI538716,	AI934035, AI799199, AI537303, AI800185,	AI783504, AL036214, AW14931	AW148320, AW087445, AA470491, AI828682,	AI349772, AI224992, AW088903, AA225339,	AI909641, AI281773, AL041150, AI690312,	AW022682, AI567351, AW074993, AW302965,	AI784252, AC006313, I48979, I89947, S68736,	AF125948, AF104032, AF090934, U42766, AC006222,	AL050149, AF090943, AL117460, AF090901,	AL050116, I48978,	AF113013,	AL110221, Y16645, AF118064, AL122050, AF177401,	AF113694, AL049452, AL133557, AF113690,	AF113019, AF113677, Y11587, AL080137, AL122123,	AF113699, AL133016, E03348, AF113689, AL049430,	AR059958, AF158248, AF146568, AC006482,	AL122121, AL137557, I49625, AL133075, Y11254,	AL050108, AL110196, AL049314, AJ000937,	AL133080, AF125949, AL050393, AL133565, X63574,	2, AL080060,	~	0, AL050146, AL137527,	3, AJ242859, AL080124,	, AF111851,	AB019565, AL049466, AL133093, A65341, AJ238278,	X96540, AC	AL122110, AL117583	X82434, AL117585, AL133113, X65873, AL137521,	AL137550	AF079765, U91329, A58524, A58523, AL049283,	E07108, AF087943, E02349	AA452295, AI700341, AA039713, AW274555,
																																Preferably excluded from the
																																877029
					_				_	_																						HOSOZ37
														·																		1719

			present invention are one or more	, AI684403, AI040232,
			polynucleotides comprising a	AA039/12, A1932286,
			nucleotide sequence described by	AA582100, AW020316,
			the general formula of a-b, where a	AA886794, AI492312, AI492311, AL034350
			is any integer between 1 to 792 of	
			SEQ ID NO:1719, b is an integer of	
_			15 to 806, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
		_	NO:1719, and where b is greater	
			than or equal to a + 14.	
1720	HCROD37	0.0278	Preferably excluded from the	
-			present invention are one or more	
			polynucleotides comprising a	
			nucleotide sequence described by	
			the general formula of a-b, where a	
			is any integer between 1 to 491 of	
			:1720, b is an	
			15 to 505, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1720, and where b is greater	
			than or equal to a + 14.	
1721	H2LAF20	877031	Preferably excluded from the	AI474074, AA313945, AW382674, AI475856, D81026,
			present invention are one or more	D59619, D80210,
			polynucleotides comprising a	D80219,
			nucleotide sequence described by	D80195, D80212, D59859, AW377671, D51799,
			the general formula of a-b, where a	D80253, D80164, D80251, D58283, D80022, D80248,
			is any integer between 1 to 665 of	D50979, D80193, D80188, C14331, D80391, D59787,
			SEQ ID NO:1721, b is an integer of	
			15 to 679, where both a and b	D57483, D80366, D80196, D59889, C15076, D80024,
			correspond to the positions of	AA305578,
			nucleotide residues shown in SEQ ID	D51022, D50995, AA514186, D80241, D80045,
			NO:1721, and where b is greater	D80378, AW177440, C14014, AA514188, C14429,
			than or equal to a + 14.	93, AW360811, D59373, T03269, '
				C75259, AW179328, C14077, AW375405, C05695,

	3
	AW360817, AW177501, AW375406, AW177511, D80439,
	AW378534, D80302, AW179332, AW377672, AW179023,
	AW178905, D80134, AW178762, D58253, D51250,
	AW178775, AW352171
	AW177731, D80247,
	AW179019, AW179024, D59627, D80258, AW352158,
	AW177505, AW352117, AW178906, AW176467,
	AW360841,
	AI910186,
	AW178908, AW178754, AW179018, AW352174, F13647,
	W179004, D58246, D58101, AW179012,
	AW178911, AW177722, AW352163, D80064, D59653,
	Z21582, AW360834, AW178983, D81111, AW178781,
	1378540,
	AW352120, T02974, C14975, H67854, H67866,
	C14298, C03092, AA809122, AW177508, AI525923,
	C14407, T03116, D51221, AI525917, D80228,
	3, AW177497, D59317, AI557774, D5
	C14973, AW177734, AIS57751, AIS2S92
	50981, AA514184, AI525215
	D60010, C14957,
	5, D59551, D60214, AI525227
	2, D52291, AW378542, AI5259
	C16955,
-	A62298, AR018138, A84916, A62300, AJ132110,
	AR008443, AB002449, AR025207, IS0126, IS0132,

				ISO128, ISO133, AR066488, AR016514, AR060138, A45456, A26615, AR052274, Y09669, A43192, A43190, AR018669, AR066490, A30438, AR066487,
				AR016691, AR016690, U46128, AB012117, I18367, I14842, AR054175, D50010, Y17187, X68127, AR008277 AR008281 A63261 A85396 D88507
				A44171, A85477, AR008408,
				A86/92, AR0628/2, A/086/, X93549, D13509, A64136, A68321, AR060133, I79511, U79457,
				AF123263, AR032065, AR008382
1722	HCROD15	877032	Preferably excluded from the	
			present invention are one or more	
			polynucleotides comprising a	
			nucleotide sequence described by	
			the general formula of a-b, where a	
			is any integer between 1 to 605 of	
			SEQ ID NO:1722, b is an integer of	
			15 to 619, where both a and b	
			correspond to the positions of	
			'O	
			Д	
			than or equal to a + 14.	
1723	HS2SG18	877034	Preferably excluded from the	AA307890
			present invention are one or more	
			polynucleotides comprising a	
			nucleotide sequence described by	
			is any integer between 1 to 838 of	•
			SEQ ID NO:1723, b is an integer of	
			15 to 852, where both a and b	
		_	correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1723, and where b is greater	
			than or equal to a + 14.	
1724	HMCHWI	180778	Preferably excluded from the	, AA307645, AL137945, R78416,
	2		present invention are one or more	AA699829, AA130430, R23973, AA204937, T58303,

	polynucleotides comprising a	AA205080, AI581369, AA130456, H03662, R77222,
	nucleotide sequence described by	965, AA1345
	the general formula of a-b, where a	AA133084, AI733757, AA088546, AA553526,
	is any integer between 1 to 683 of	AA843823, AW392930, AI522161, AA055592, R66492,
	SEQ ID NO:1724, b is an integer of	AI820789, AI
	15 to 697, where both a and b	I499378, AA151971,
	correspond to the positions of	AA479719, AA100721
	$\boldsymbol{\sigma}$	AL050348, AL035419, AC005276, AL121782,
	NO:1724, and where b is greater	AL080316, AC007617, AC010168, AC008069,
	than or equal to a + 14.	AC000064, AC002984, AB020874, AC007401,
		AC007566, AC005150, AC005145, AC007022,
		AL035067, AC000114, AC007685, AC005549,
		AC007207, AC006146, AL031767, AC008072,
		, AF130342, AL035408
		AC007681, AC008134, Z92543, AJ133269, AC005386,
		AL049546, AC004998, D11078, AC004986, AL035698,
_		
_		AC005090, AC004514, AC005837, AC003013,
		AL009031, AC007463, AC009946, AC006364,
-		, AC00541
		Z82210, AL139054, AL022068, AL121718, AC007381,
		, AF118808
		, AC005036,
_		, AL133224,
		, AF053936,
		, AC004072,
		, AB020871, AL021327,
		AL021940,
		AC005234,
		AF049895, AC006382, Z95327, AL031073, AL117327,
		AC005392, AC007001, AL035610, AC002384, U95626,
		AC007785, 299495, AL109809, AF149773, AF068862,
		AC005154,
		AC005531,
		AL109967, AC004617, AP000230, AP000144,

AC004858, AC007276, AC011604, AC005723, AL132987, AF011889, AP000155, AL050325, AC006582, AC004924, D87055, AC004472, AP AC005686, AL13371, AF026249, AL02330, AF110315, AF108841, Z83818, AL034350, D1 AF064074, AF064073, AI698170, AI346506,						1			
HWLVSS2 877043 Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 454 of SEQ ID NO:1725, b is an integer of 15 to 468, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1725, and where b is greater than or equal to a + 14. HCRPG56 877044 Preferably excluded from the polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 468 of SEQ ID No:1726, b is an integer of 15 to 482, where both a and b correspond to the positions of 15 to 482, where both a and b correspond to the positions of 15 to 482, where both a and b correspond to the positions of 15 to 482, where both a and b correspond to the positions of 15 to 482, where both a and b correspond to the positions of 15 to 482, where both a and b correspond to the positions of 15 to 482, where both a and b correspond to the positions of 15 to 482, where both a and b correspond to the positions of 15 to 482, where both a and b correspond to the positions of 15 to 482, where both a and b correspond to the positions of 15 to 482, where both a and b correspond to the positions of 15 to 482, where both a and b correspond to a + 14.					AL022318, AC004510,		.C007276, .C005723,	AF109907, AL079352,	
HWLVSS2 877043 Preferably excluded from the present invention are one or more polynucleotides comprising a mucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 454 of SEQ ID NO:1725, b is an integer of 15 to 468, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1725, and where b is greater than or equal to a + 14. HCRPG56 877044 Preferably excluded from the polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a six any integer between 1 to 468 of SEQ ID NO:1726, b is an integer of nucleotide residues shown in SEQ ID NO:1725, and where both a and b correspond to the positions of 15 to 482, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1726, and where b is greater than or equal to a + 14. HTAHC75 877046 Preferably excluded from the AI916318,					AC002326,		F011889,	AL049544,	
HWLVSS2 877043 Preferably excluded from the polynucleotide sequence described by AF005632, AF005					AP000013,		L050325,	AC007182,	
HWLVSS2 877043 Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 454 of SEQ ID NO:1725, b is an integer of 15 to 468, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1725, and where b is greater than or equal to a + 14. HCRPGS6 877044 Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 468 of SEQ ID NO:1726, b is an integer of 15 to 482, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1726, and where b is greater than or equal to a + 14. HTAHCTS 877046 Preferably excluded from the Al916318.					AL035690,	AC006582, A	C004924,	,	276735,
HWLVSS2 877043 Preferably excluded from the present invention are one or more polynuclectides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 454 of SEQ ID NO:1725, b is an integer of 15 to 468, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1725, and where b is greater than or equal to a + 14. HCRPGS6 877044 Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 468 of SEQ ID NO:1726, b is an integer of 15 to 482, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1726, and where b is greater than or equal to a + 14. HTAHCTS 877046 Preferably excluded from the AB1916318,					AC006459,	D87055, AC0	04472, AF	2000501, AC005002	5002,
HWLVSS2 877043 Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 454 of SEQ ID NO:1725, b is an integer of 15 to 468, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1725, and where b is greater than or equal to a + 14. HCRPG56 877044 Preferably excluded from the present invention are one or more present invention are one or more present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 468 of SEQ ID NO:1726, b is an integer of 15 to 482, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1726, and where b is greater than or equal to a + 14. HTAHCCS 877046 Preferably excluded from the A1916318.					AF205592,		L133371,	AF026248,	
HWLVSS2 877043 Preferably excluded from the present invention are one or more polynuclectides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 454 of SEQ ID NO:1725, b is an integer of 15 to 468, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1725, and where b is greater than or equal to a + 14. HCRPG56 877044 Preferably excluded from the present invention are one or more present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 468 of SEQ ID NO:1726, b is an integer of 15 to 482, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1726, and where b is greater than or equal to a + 14. HTAHCT5 877046 Preferably excluded from the A1916318,					AF026254,	AF026249, A			
HWLVS52 877043 Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotides comprising a nucleotides comprising a nucleotides comprising a nucleotides comprising a sis any integer between 1 to 454 of SEQ ID NO:1725, b is an integer of 15 to 468, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1725, and where b is greater than or equal to a + 14. HCRPG56 877044 Preferably excluded from the polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 468 of 15 any integer both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1726, and where b is greater than or equal to a + 14. HTAHC75 877046 Preferably excluded from the A1916318.					AF108842,	AF110315, A		AF108843,	
HWLVS52 877043 Preferably excluded from the present invention are one or more polynuclectides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 454 of SEQ ID NO:1725, b is an integer of 15 to 468, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1725, and where b is greater than or equal to a + 14. HCRPG56 877044 Preferably excluded from the polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 468 of SEQ ID NO:1726, b is an integer of 15 to 482, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1726, and where b is greater than or equal to a + 14. HTAHC75 877046 Preferably excluded from the AI916318, AI698170, AI346506,					AC007280,	Z83818, AL0	34350, D1	10083, AC0030	
HWLVSS2 877043 Preferably excluded from the present invention are one or more polynucleotides comprisined by the general formula of a-b, where a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 454 of SEQ ID NO:1725, b is an integer of 15 to 468, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1725, and where b is greater than or equal to a + 14. HCRPGS6 877044 Preferably excluded from the polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 468 of SEQ ID NO:1726, b is an integer of 15 to 482, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1726, and where b is greater than or equal to a + 14. HTAHCTS 877046 Preferably excluded from the A1916318, A1698170, A1346506.					AC005632,	4	F064073,	AC007556, AC	AC004889
present invention are one or more polynuclectides comprising a nuclectide sequence described by the general formula of a-b, where a is any integer between 1 to 454 of SEQ ID NO:1725, b is an integer of 15 to 468, where both a and b correspond to the positions of nuclectide residues shown in SEQ ID NO:1725, and where bis greater than or equal to a + 14. HCRPGS6 877044 Preferably excluded from the polynuclectides comprising a nuclectide sequence described by the general formula of a-b, where a is any integer between 1 to 468 of SEQ ID NO:1726, b is an integer of 15 to 482, where both a and b correspond to the positions of nuclectide residues shown in SEQ ID NO:1726, and where b is greater than or equal to a + 14. HTAHCTS 877046 Preferably excluded from the AI916318, AI698170, AI346506.	1725	HWLVS52	877043	Preferably excluded from the					
polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 454 of SEQ ID NO:1725, b is an integer of 15 to 468, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1725, and where b is greater than or equal to a + 14. HCRPGS6 877044 Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 468 of SEQ ID NO:1726, b is an integer of 15 to 482, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1726, and where b is greater than or equal to a + 14. HTAHCT5 877046 Preferably excluded from the Al916318, Al698170, Al346506.				Ä					
nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 454 of SEQ ID NO:1725, b is an integer of 15 to 468, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1725, and where b is greater than or equal to a + 14. HCRPG56 877044 Preferably excluded from the polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 468 of SEQ ID NO:1726, b is an integer of 15 to 482, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1726, and where b is greater than or equal to a + 14. HTAHC75 877046 Preferably excluded from the				polynucleotides comprising a					
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is any integer between 1 to 454 of SEQ ID NO:1725, b is an integer of 15 to 468, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1725, and where b is greater than or equal to a + 14. HCRPG56 877044 Preferably excluded from the polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 468 of SEQ ID NO:1726, b is an integer of 15 to 482, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1726, and where b is greater than or equal to a + 14. HTAHC75 877046 Preferably excluded from the A1916318, A1698170, A1346506,									
SEQ ID NO:1725, b is an integer of 15 to 468, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1725, and where b is greater than or equal to a + 14. HCRPG56 877044 Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 468 of SEQ ID NO:1726, b is an integer of 15 to 482, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1726, and where b is greater than or equal to a + 14. HTAHC75 877046 Preferably excluded from the				is any integer between 1 to 454 of					
15 to 468, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1725, and where b is greater than or equal to a + 14. HCRPG56 877044 Preferably excluded from the polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 468 of SEQ ID NO:1726, b is an integer of 15 to 482, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1726, and where b is greater than or equal to a + 14. HTAHC75 877046 Preferably excluded from the A1916318, A1698170, A1346506,				:1725, b is an integer					
nucleotide residues shown in SEQ ID NO:1725, and where b is greater than or equal to a + 14. HCRPGS6 877044 Preferably excluded from the polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 468 of SEQ ID NO:1726, b is an integer of 15 to 482, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1726, and where b is greater than or equal to a + 14. HTAHC75 877046 Preferably excluded from the AI916318, AI698170, AI346506,				, where both a					
NO:1725, and where b is greater than or equal to a + 14. HCRPG56 877044 Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 468 of SEQ ID NO:1726, b is an integer of correspond to the positions of nucleotide residues shown in SEQ ID NO:1726, and where b is greater than or equal to a + 14. HTAHC75 877046 Preferably excluded from the A1916318, A1698170, A1346506,				ons					
HCRPG56 877044 Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 468 of SEQ ID NO:1726, b is an integer of 15 to 482, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1726, and where b is greater than or equal to a + 14. HTAHC75 877046 Preferably excluded from the A1916318, AI698170, AI346506,									
HCRPG56 877044 Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 468 of SEQ ID NO:1726, b is an integer of 15 to 482, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1726, and where b is greater than or equal to a + 14. HTAHC75 877046 Preferably excluded from the Al916318, Al698170, Al346506,		-		NO:1725, and where b is greater					•
HCRPG56 877044 Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 468 of SEQ ID NO:1726, b is an integer of 15 to 482, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1726, and where b is greater than or equal to a + 14. HTAHC75 877046 Preferably excluded from the AI916318, AI698170, AI346506,				equal to a + 14.					
present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 468 of SEQ ID NO:1726, b is an integer of 15 to 482, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1726, and where b is greater than or equal to a + 14. HTAHC75 877046 Preferably excluded from the AI916318, AI698170, AI346506,	1726	HCRPG56	877044		١.	١.		AC009533, AC00	AC008013
polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 468 of SEQ ID NO:1726, b is an integer of 15 to 482, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1726, and where b is greater than or equal to a + 14. HTAHC75 877046 Preferably excluded from the A1916318, A1698170, A1346506,				ΣC					
nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 468 of SEQ ID NO:1726, b is an integer of 15 to 482, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1726, and where b is greater than or equal to a + 14. HTAHC75 877046 Preferably excluded from the A1916318, A1698170, A1346506,				polynucleotides comprising a					
the general formula of a-b, where a is any integer between 1 to 468 of SEQ ID NO:1726, b is an integer of 15 to 482, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1726, and where b is greater than or equal to a + 14. HTAHC75 877046 Preferably excluded from the AI916318, AI698170, AI346506,				nucleotide sequence described by					
is any integer between 1 to 468 of SEQ ID NO:1726, b is an integer of 15 to 482, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1726, and where b is greater than or equal to a + 14. HTAHC75 877046 Preferably excluded from the AI916318, AI698170, AI346506,									
SEQ ID NO:1726, b is an integer of 15 to 482, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1726, and where b is greater than or equal to a + 14. HTAHC75 877046 Preferably excluded from the AI916318, AI698170, AI346506,				is any integer between 1 to 468 of					
15 to 482, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1726, and where b is greater than or equal to a + 14. HTAHC75 877046 Preferably excluded from the AI916318, AI698170, AI346506,									-
Correspond to the positions of NUCleotide residues shown in SEQ ID NO:1726, and where b is greater than or equal to a + 14. HTAHC75 877046 Preferably excluded from the AI916318, AI698170, AI346506,				and					
nucleotide residues shown in SEQ ID NO:1726, and where b is greater than or equal to a + 14. HTAHC75 877046 Preferably excluded from the AI916318, AI698170, AI346506,				correspond to the positions of					•
NO:1726, and where b is greater than or equal to a + 14.									
HTAHC75 877046 Preferably excluded from the AI916318, AI698170, AI346506,				and where b is					
HTAHC75 877046 Preferably excluded from the AI916318, AI698170, AI346506,				equal to a +					
	1727	HTAHC75	877046	from	AI916318,	-	1346506,	AA481006,	

			present invention are one or more	AW006462, AI808371, AI492123, AI860659,	
			polynucleotides comprising a	AW083792, AI298294, AI377296, AI299866,	_
			nucleotide sequence described by	AI143985, AI832385, T66213, AA315944, AA774467,	67,
			the general formula of a-b, where a	AA481745, AA745359, N78840, AA744416, AA03564	44,
			teger between 1 to 1883	AW236811, AI693629, AI299645, R54532, AA98735	58,
			SEQ ID NO:1727, b is an integer of	AW136153,	98,
			15 to 1897, where both a and b	AI459849, R55684, R99148, AA975345, R45317,	
			correspond to the positions of	H08045, AA992883, AI122963, AA987223, H18288,	_
			nucleotide residues shown in SEQ ID	AI681364, R55685, F09827, H46943, AW418590,	
			NO:1727, and where b is greater	1745480,	
			than or equal to a + 14.	AW192055, AA972155, R14680, F04052, AA827984,	
				F12197, H26802, T29943, AA295772, R38093,	
				AI290682, AL047550, T07816, AA355247, H07939,	_
				H69808, R38173, T85773, R54435, AA508768,	
					26,
				H51338, F11088, AA916514, T77104, R42403,	
				O, AA3005	
1728	HCRPH26	877047	Preferably excluded from the	AF118076	
			present invention are one or more		
			polynucleotides comprising a		
			nucleotide sequence described by		
			the general formula of a-b, where a		
			is any integer between 1 to 509 of		
			SEQ ID NO:1728, b is an integer of		
			15 to 523, where both a and b		
			correspond to the positions of		
			nucleotide residues shown in SEQ ID		
			NO:1728, and where b is greater		
			than or equal to a + 14.		
1729	HWLWL67	877049	Preferably excluded from the	AI620255, AI739424, AW008095, N6437	73,
			present invention are one or more	AA628778, AI827544, AI246150, AA977500,	
			polynucleotides comprising a	AA779757, AI216037, AA724806, AI143969,	
			nucleotide sequence described by	AA953515,	
			the general formula of a-b, where a	AA971965, AA010881, AI352432, AA410372,	
			is any integer between 1 to 204 of	AW082274, AA129683, AI699673, AI807260,	ļ

			SEO ID NO.1729 bis an integer of	A1375466 A1633645 AA588195 AA670218.
			15 to 218, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEO ID	
			NO:1729, and where b is greater	
			than or equal to a + 14.	
1730	HOSDU39	877050	Preferably excluded from the	
			present invention are one or more	
			polynucleotides comprising a	
			nucleotide sequence described by	
	_		the general formula of a-b, where a	
			is any integer between 1 to 566 of	
			SEQ ID NO:1730, b is an integer of	
			15 to 580, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1730, and where b is greater	
			equal to a + 14.	
1731	HCROS68	877051	Preferably excluded from the	AI940522, AC007688
			present invention are one or more	
			polynucleotides comprising a	
		_	nucleotide sequence described by	
			the general formula of a-b, where a	
			is any integer between 1 to 623 of	
			SEQ ID NO:1731, b is an integer of	
			15 to 637, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1731, and where b is greater	
		·	than or equal to a + 14.	
1732	HWLRT47	877052	Preferably excluded from the	AA676521
			present invention are one or more	
			polynucleotides comprising a	
			nucleotide sequence described by	
			the general formula of a-b, where a	
			is any integer between 1 to 409 of	

			90 TO NO. 1722 h . a. h . a. CET OED	
			correspond to the positions of	
	•		nucleotide residues shown in SEQ ID	
			NO:1732, and where b is greater	
			than or equal to a + 14;	
1733	HCRPN44	877056	Preferably excluded from the	AI814630, AI659745, AI337185, AI476215,
			present invention are one or more	AW014950, W90223, AI683180, AI040605, AI052156,
			polynucleotides comprising a	AW419172, N20981, N92247, AI583402, N51526,
			nucleotide sequence described by	H64280, H64281, H21597, AW117231, W37142,
			the general formula of a-b, where a	W47567, H65040, Z40718, H65039, W86558, W90127,
			is any integer between 1 to 1267 of	AI572195
-			SEQ ID NO: 1733, b is an integer of	R16990, AA002167,
			15 to 1281, where both a and b	
			correspond to the positions of	H66212, H66857, N30250, W15238, W15419,
			nucleotide residues shown in SEQ ID	AA024406, AA076483, AA099706, AA513421,
			NO:1733, and where b is greater	AA535580, AA593084, AA593075, AA639881,
			than or equal to a + 14.	
				AA705190, AA775052, AA854917, AI085171,
				AA952891, AA952941, AI307637, AI348056,
				AI203039, AI380800, AI473584, AI571026,
				AI424140, AI219098, AI659256, AI636785, AI338942
1734	HCRPD33	877057	Preferably excluded from the	AI167356, AL049670, AL021397
			polynucleotides comprising a	
-			nucleotide sequence described by	
			the general formula of a-b, where a	
			is any integer between 1 to 261 of	
			SEQ ID NO:1734, b is an integer of	
			15 to 275, where both a and b	
			correspond to the positions of	
	-		nucleotide residues shown in SEQ ID	
			NO:1734, and where b is greater	
			than or equal to a + 14.	
1735	HCRPE57	877058	Preferably excluded from the	AA989345, AI624083, D61985, N67616
			present invention are one or more	

			otides comp	AW303456, AA456790, AI051183, AW152159,
				AA130046, R/9256, AW439608, H22118, AA134040,
			the general formula of a-b, where a	
			is any integer between 1 to 778 of	T79540, T97240, T97241, R51919, AW079574,
		-	SEQ ID NO:1738, b is an integer of	C00464, AI699839, AI689564, AL046171, AI702873,
			15 to 792, where both a and b	R79157, AI905847, AA129873, AA356980, AA351418,
			correspond to the positions of	T09084, AW248101, AI929724, AI815427, W27745,
			nucleotide residues shown in SEQ ID	D85131, M94046, AB017335, M93339, U33819
			NO:1738, and where b is greater	
			than or equal to a + 14.	
1739	HWLVE77	877066	Preferably excluded from the	N53758
			present invention are one or more	
			polynucleotides comprising a	
			nucleotide sequence described by	
			the general formula of a-b, where a	
			is any integer between 1 to 454 of	
			SEQ ID NO:1739, b is an integer of	
			15 to 468, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1739, and where b is greater	
			than or equal to a + 14.	
1740	HCROJ64	877067	Preferably excluded from the	
			present invention are one or more	
			polynucleotides comprising a	
			nucleotide sequence described by	
			the general formula of a-b, where a	
			is any integer between 1 to 93 of	
			SEQ ID NO:1740, b is an integer of	
			15 to 107, where both a and b	
		_	correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1740, and where b is greater	
			than or equal to a + 14.	
1741	нмгомо	877068	Preferably excluded from the	
	5		present invention are one or more	

		AW392670, U46347, Z99396, AW363220, AW372827, AL119484, AL119457, AL119324, AL119324, AL119355, AL119341, AL119483, AL119522, AL043003, U46351, U46349, AL119444, U46350, U46341, AL119396, AL119496, AL119418, AL043041, AL119396, AL119418, AL043053, AL042614, AL119399, AL1134538, U46345, AL042450, AL042544, AL043019, AL043029, AL119464, AC015853, AR060234, A81671, AB026436, AR054110, AR069079
		U46347, Z99396, AW AL119484, AL119457 AL119341, AL119483 AL146350, U46351, U46350, U46341, AL AL13453, AL134528 AL119418, AL043033, AL042984, AL042965, AL042984, AL043049, AL042551, AL043029, AL042551, AL043147, AC015853, AR060234, AR054110, AR069079
		AW392670, AL119363, AL119355, AL119522, AL119444, AL119446, AL119446, AL11946, AL119399, AL119399, AL119464, AL119464, AL119464,
	AC004540	AA906013, AL119319, AL119391, AL119439, AL119439, AL119439, AL119335, AL037205, AL134153, AL042975, AL042542, AL119304, AL119304,
polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 471 of SEQ ID NO:1741, b is an integer of 15 to 485, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1741, and where b is greater than or equal to a + 14.	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 398 of SEQ ID NO:1742, b is an integer of 15 to 412, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1742, and where b is greater than or equal to a + 14.	preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 380 of SEQ ID NO:1743, b is an integer of 15 to 394, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1743, and where b is greater than or equal to a + 14.
	877069	877070
	HCRPW24	HOCTA26
	1742	1743

1744	HBKDB96	877071	Preferably excluded from the	200712002	713500AC	ONSCOOL	710070	
			present invention are one or more	AA931328,	AI392998,	AI287567,	A1004013, A1493596,	
			polynucleotides comprising a	AI278360,	H16208, AM	1375190, HS	1009, AW375161	5161,
			nucleotide sequence described by	AW375154,	AW375158,	H90897, H16	5209, AW375149	5149,
			eral formula of a-b,	AW418706,	AW385279			
				:		•		
			SEQ ID NO:1744, b is an integer of					_
			15 to 953, where both a and b					
			correspond to the positions of					
			nucleotide residues shown in SEQ ID					
,			NO:1744, and where b is greater	-				_
			than or equal to a + 14.					
1745	HCRPE30	877073	Preferably excluded from the	AB014604,	AC003093			
			present invention are one or more					
_			polyhucleotides comprising a					
			nucleotide sequence described by					-
			the general formula of a-b, where a					
-			is any integer between 1 to 378 of					
			SEQ ID NO:1745, b is an integer of					
_			15 to 392, where both a and b					
			correspond to the positions of					-
			de residues					
			NO:1745, and where b is greater					
			than or equal to a + 14.					
1746	HKGAW02	877075	Preferably excluded from the	AA935168,	AA398801,	AL119484, A	AL134524, 1	AL119418
			present invention are one or more					
_			polynucleotides comprising a					
	-		nucleotide sequence described by					
			the general formula of a-b, where a					
			is any integer between 1 to 519 of					
			SEQ ID NO:1746, b is an integer of					
	_		15 to 533, where both a and b					-
			correspond to the positions of					
		_	·O					
			NO:1746, and where b is greater					
			than or equal to a + 14.		İ			_

AI434772		R13359, H08041, AF010245, AW156983, H29189, Z46132, T16980, AI879608, AW402188, AA348764, R34542, R61072, H23510, AA436740, N36381, AI929579, AI879056, AI816318, AL137450
Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 237 of SEQ ID NO:1747, b is an integer of 15 to 251, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1747, and where b is greater than or equal to a + 14.	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 341 of SEQ ID NO:1748, b is an integer of 15 to 355, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1748, and where b is greater than or equal to a + 14.	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 818 of SEQ ID NO:1749, b is an integer of 15 to 832, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1749, and where b is greater than or equal to a + 14.
877079	877080	877083
нсосряз	НОСТD62	HE8PC46
1747	1748	1749

1750	HWI OMS	77087	Dreferably excluded from the	AMAGGGG	ALGACATA	LEGISTRE SISTER	ייטראנא
2			dry characa rroll cile	AND 02003,	A101/014,	WAYOVOTO'	AA/019/1,
	າ			AA465292, AA204693	AA204693		
			polynucleotides comprising a				
			nucleotide sequence described by				
			the general formula of a-b, where a				
			is any integer between 1 to 470 of				
			SEQ ID NO:1750, b is an integer of				
			15 to 484, where both a and b				
			correspond to the positions of				
			nucleotide residues shown in SEQ ID				
			NO:1750, and where b is greater				
			than or equal to a + 14.				
1751	HTLGE26	877088	Preferably excluded from the	AI285916,	AI025315,	AP000553,	AC009516
			present invention are one or more				
			polynucleotides comprising a				
			nucleotide sequence described by				
			the general formula of a-b, where a				
			is any integer between 1 to 758 of				
			SEQ ID NO:1751, b is an integer of				
			15 to 772, where both a and b				
			correspond to the positions of				
			nucleotide residues shown in SEQ ID				
			NO:1751, and where b is greater				
			than or equal to a + 14.				
1752	HCFDE85	877092	Preferably excluded from the				
			present invention are one or more				
			polynucleotides comprising a				
			nucleotide sequence described by				
			the general formula of a-b, where a				
			is any integer between 1 to 370 of				
			SEQ ID NO:1752, b is an integer of				
			15 to 384, where both a and b				
			correspond to the positions of				
			nucleotide residues shown in SEQ ID				
			NO:1752, and where b is greater				
			than or equal to a + 14.				

1753	HFEAH85	877093	Preferably excluded from the	AI950320, AA340023
			present invention are one or more	
			polynucleotides comprising a	
			nucleotide sequence described by	
			the general formula of a-b, where a	
			is any integer between 1 to 208 of	
			SEQ ID NO:1753, b is an integer of	
			15 to 222, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1753, and where b is greater	
			than or equal to a + 14.	
1754	HE8QT45	877094	Preferably excluded from the	AI052389, AI761986, AW057796, AI656751,
	,		present invention are one or more	AW152082, AI126366, AI125599, AA452171,
			polynucleotides comprising a	AI687797, AW023851, AA406351, AI431689,
			nucleotide sequence described by	AA778840, AA993437, AI128983, AA565214,
			the general formula of a-b, where a	AI693581, AI254753, AI285759, AW020705,
			is any integer between 1 to 636 of	AI762885, N92604, AI193254, AI003334, C16412,
				C16192, AA226919, AA479128, AI536542, H08761,
			15 to 650, where both a and b	AA706764, R85597, T10616, AI933471, AI250282,
			correspond to the positions of	AW160916, AI440238, AW151132, AI372041,
			nucleotide residues shown in SEQ ID	AL040011, AA731417, AA806605, AA641818,
			NO:1754, and where b is greater	AW194014, AA938181, AI932739, AW020164,
			than or equal to a + 14.	AI345688, AI813538, AA829402, AI431507,
				, AW080157, AI963101,
				, AW167340, AW151974
				н
				AA609644, AI627339, AI499057, AI690813,
				AI581053, AI866469, AI955441, AW021373,
				AA282824, AI799313, AI609409, AA810226,
				AI918449, AI699029, AW189548, AW058304,
				AI828676, AI659041, AI918809, AA065052,
				AL134828, C21335, AI357644, AI348821, AI590043,
				AI866770, AI399759, AI636507, AA767924,
				AA814517, AI289791, AI421662, AW082532,
				AA761557, AA743474, AA836665, AI628850,

6, AW088546, AI590755, W48671,	, AI241923, AL079963, AI44	, AI884574,	AI581033, AW148544, AW079996, AA811736,	8, AW078818, AW409793,	AW002727, AI859991, AI688381, AW406745,	AW021717, AW196720, AI915291, AW152182,	AI950729, AI472487, AW023072, AI921915,	AI582932, AI609191, AI872423, AI619820,	AI434731, AI524179, AI800370, AI521560,	AI889189, AW075382, N52016, AW089844, AI648494,	AI678623, AI273886, AW104141, AW029457,	AL022334, AR050959, S75997, AF100931, AF141289,	AF183393, A18777, AL133619, AF039138, AF039137,	A08910, A08909, AF103804, AL110269, AB020777,	AR038854	5, AF000167, A76337,	AF067790, AL137537, AL050155, AR053103, I48978,	X55761, AF036941, Y13653, I89947, I33392,	AC010077, AF026816, I80062, X83544, I22020,	X99270, AF044323, X66366,	X01775, A18788, X80340, AC006288, AL133565,	AL137479, A60092, A60094, AF031572, AC004383,	249216,	I77092, D5	A65340, AL	A77035, AL122104, AL137271, E03168, AF184965,	X93328, AL	AF038847, AL137554, AF043493, AL110158, AF042090	W79030, AC005486				
													-													_			Preferably excluded from the	present invention are one or more	ides comp	eotide sequence describ	the general formula of a-b, where a is any integer between 1 to 546 of
																													877095				
				-			·	-																					HWLQL84			•	
												-																	1755				

			1 2 5		Г
			O:1/33, D is all illeger		
			15 to 560, where both a and b		
			nucleotide residues shown in SEQ ID		
			NO:1755, and where b is greater		
			than or equal to a + 14.		\neg
1756	HCQCP82	877096	Preferably excluded from the	AA193032	
			present invention are one or more		
_			polynucleotides comprising a		
			nucleotide sequence described by		
			the general formula of a-b, where a		
			is any integer between 1 to 275 of		
			SEQ ID NO:1756, b is an integer of		
			15 to 289, where both a and b		
			correspond to the positions of		
			nucleotide residues shown in SEO ID		
			disc. 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1		
			than or equal to a + 14.		П
1757	HCRMW8	877097		AI902587, AL110283	
	0		present invention are one or more		
			polynucleotides comprising a		
			nucleotide sequence described by		
			the general formula of a-b, where a		
			is any integer between 1 to 476 of		
			SEQ ID NO:1757, b is an integer of		
			15 to 490, where both a and b		
			correspond to the positions of		
			nucleotide residues shown in SEQ ID		
			NO:1757, and where b is greater		
			than or equal to a + 14.		
1758	HSIGL73	877098	Preferably excluded from the		
			present invention are one or more	AA248302, AI537677, AI345416, AI345612,	
			polynucleotides comprising a	AI345415, AL134830, AI802542, AW051258,	
			de s	, AI677796,	
			the general formula of a-b, where a	AI619502, AW198090, AI433157, AI702073,	_
			is any integer between 1 to 841 of	AI633125, AI334445, AW163464, AI254727,	

				-					_											_					-										
98	AI521012,	AW148716,	AI554245,	AI611738,	AW073865,	AI954183,	AW300889,	AL040243,	AI933589,	AL039086,	AI612885,	AI697324,	AW169653,	AI500523,	AI521560,	AW104827,	A1475371,	AL036403,	AA427700,	AL036631,	AW301409,	AW150578,	AL045500,	AI815855,	AW020561,	AA572758,	AI587114,	AI815237,	AW268220,	AL120853,	AI439745,	AW087207,	AI249323,	AW087445,	AW020693.
AI499285,	AIS64719,	AL036736,	AW090071,	AI358701,	AI536638,	AI687362,	AI887659,	AI921248,	AI873644,	AI620284,	AI919534,	AW129659,	AL038069,	AI801325,	AI926790,	AW023590,	AI491852,	AW190194,	AI283760,	AI699865,	AI207510,	AL036980,	AW080402,	AL043293,	AI702068,	AI559296,	AW073270,	AI469532,	AI468872,	AW166970,	AIS00706,	AI590120,	AI862144,	AI696398,	AW081298,
	AI886753,	AL119863,	AI340603,	AL046200,	AI445025,	AA640779,	AI571909,	AW117746,	AI627360,	AI783504,	AI637584,	AW163823,	AI670009,	AI612913,	AL037454,	AW090013,	AI348897,	AI520862,	AW148363,	AL036274,	AI524671,	AI886124,	AI440239,	AW075667,	AL036396,	AI866770,	AW193530,	AI312428,	AI536685,	AI340519,	AI932794,	AI648509,	AI433976,	AI934259,	AA470491,
സ	AI863241,	AW026882,	AW161579,	AW160916,	AI284131,	AI636588,	AW300782,	AI500077,	AI632408,	AI682743,	AL120307,	AI815232,	AI284517,	AW104724,	AI446373,	AI500662,	AI890833,	AI627988,	AI567128,	AI284484,	AI798456,	AI812107,	AI679504,	AW118518,	AW148408,	AL038605,	AL040241,	AI610690,	AI866801,	AI805603,	AI349645,	AW089572,	AL110306,	AI280747,	AI929108,
0:1758, b is an inte		to the po	nucleotide residues shown in SEQ ID	•	than or equal to a + 14.																														
				_																													•		

	_		AWIOSKOI	AW193911	AT620866	AT306613.	_
			1 1 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2	1100011			
			AI274541,	AI609375,	A156/612,	AW022808,	
			AL036802,	AI270055,	AI174394,	AI554186,	
			AW129916,	AI613270,	AI633330,	AI874166,	
	_		AI625079,	AI683585,	AL047763,	AW132056,	
			AW169527,	AI335426,	AI348777,	AI270099,	
	.,.		AI862139,	AI355827,	AI475394,	AI285448,	
			AI687065,	AI686576,	AA806720,	AI871697,	
			AW403717,	AI682971,	AL036361,	N33175, AI889376,	.,`
			AI923989,	AW152459,	AI636585,	AI439717,	
			AL119791,	AI635461,	AI433384,	AI923370,	
	_		AI345131,	AIS91075,	AI567351,	AW074993,	
			AW302965,	AI431424,	AI349614,	AW193134,	
			AI343112,	AI954422,	AI434468,	AI499986,	
	_		AW268083,	AI572787,	AW268253,	AI537515,	
			AI281772,	AL045266,	AI254731,	AI349598,	
			AI934011,	AI312152,	AI872545,	AI570807,	
			A1686817,	AI247293,	AL041772,	AI345735,	
			AI819326,	AW078839,	AI539771,	AW075084,	
			AI818977,	AI784252,	Z83839, L29339,	29339, AF042090,	
			AC004057,	AL032822,	AC004470,	AL080239,	
			AC018767,	AC006197,	AC004554,	AC004808,	
			AC006313,	AC002454,	AF090900,	AL133560,	•
			AF090934,	AL137271,	I48978, I	I48978, I89947, A08916,	
			AL133557,	AL117460,	AL049382,	AJ000937,	
			AL049314,	AF111851,	AC002480,	A08913	
1759 HHEYT40	877099	Preferably excluded from the	AA313905,	AW392670,	AL119319,	9	
		present invention are one or more	AW372827,	U46349, A	AL119399, AL119363,	AL13451	, ,
		polynucleotides comprising a	AL119443,	AW363220,	AW363220, AW384394, U46346,		
		nucleotide sequence described by	AL119497,	U46347, A		L119335, AL134528	
		al	U46351, AL042850,		AL119457, A	AL119522, AL13492(. ,
		teger between	AL119484,	AL119391,	AL119324, AL119444	, 29939	6,
		ID NO	AL119355,	AL119483,	U46345, A	U46345, AL134538, AL11943	,
			AL043037,	AL042970,	AL037205,	AB026436, A81671	
		correspond to the positions of	AR054110,	AR060234,	AR066494		
		Q.					

			NO:1759, and where b is greater	
			than or equal to a + 14.	
1760	нронозі	877101	Preferably excluded from the	AW405179, AA278430, AI951459, AW130135,
			present invention are one or more	AA437355, AA427621, AW183077, AW044380,
-		_	polynucleotides comprising a	AI038334, AI540554, AI224500, AA256905,
			nucleotide sequence described by	AW440059, AA702920, AI269240, AA662464,
			the general formula of a-b, where a	AA129087, AI042498, AW401902, AI865421,
			is any integer between 1 to 2712 of	AA129086, AI023674, AA670374, U51141, AI355031,
			SEQ ID NO:1760, b is an integer of	
		_	15 to 2726, where both a and b	AA278961, AI286001, AW237708, AA512902, R16374,
			correspond to the positions of	AI000189, AA872607, Z39825, AW338997
			nucleotide residues shown in SEQ ID	
			NO:1760, and where b is greater	
			than or equal to a + 14.	
19/1	HODGR31	877104	Preferably excluded from the	AI701474, AI141563, AA805242, AW151887,
			present invention are one or more	AW172894, AI342500, N26482, AI990393, AW275998,
			polynucleotides comprising a	AL120029, AI367540, AA905238, AA767195,
			nucleotide sequence described by	AA633403, N25228, AA811725, Z39323, N29704,
			the general formula of a-b, where a	H17935, W05575, N70530, AA766858, AL118631,
			is any integer between 1 to 1019 of	N98948, AI701701, N66665, AA737077, AB007917
			SEQ ID NO:1761, b is an integer of	
			15 to 1033, where both a and b	
			correspond to the positions of	
			Ö	
			NO:1761, and where b is greater	
			than or equal to a + 14.	
1762	HWLWB9	877105	Preferably excluded from the	AA167624, AA688144, AA016314, AI499580,
	2		present invention are one or more	AI925014, AA808419, AI081193, AA194836,
			polynucleotides comprising a	AA125835, AW419229, AA252083, AA461554,
			nucleotide sequence described by	
			the general formula of a-b, where a	AA687098, W33019, AA876407, AW007949, F34751,
			is any integer between 1 to 607 of	AA492322, AA908820, R37941, T23517, AA844143,
			Π.	N73484, AA488062
			15 to 621, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	

			NO:1762, and where b is greater	
			than or equal to a + 14.	
1763	HWLRD79	877106	Preferably excluded from the	H51960, AA393998, AI
			present invention are one or more	AI017517, AI819082, AW088106, AW264111,
			polynucleotides comprising a	AI446796, AA767844, AI538119, AI583021,
	•		nucleotide sequence described by	AW151792, AW168958, AI252808, T79312, AA429868,
			the general formula of a-b, where a	AI358328, AI039023,
			is any integer between 1 to 722 of	
			SEQ ID NO:1763, b is an integer of	N64391, AI275601, AA437374, AW003543, H93076,
			15 to 736, where both a and b	AI962621, AI148567, AA904883, AW194543, F01936,
			correspond to the positions of	AI674414, AI419876, AI339747, AW299722, C00822,
			nucleotide residues shown in SEQ ID	AA661775, T27646, AI473622, AI473612, AL042432,
			NO:1763, and where b is greater	AA775934, AA700143, X63546, I76205, AJ012755
			than or equal to a + 14.	
1764	HWLOW7	877110	Preferably excluded from the	
	2		present invention are one or more	AI128724, AI990335, AA456529, AI655816, H39555,
			polynucleotides comprising a	AI479968, AI283132, AI926934, AA534329,
			nucleotide sequence described by	AA019380, AI961572, AA011475, AI089295,
			the general formula of a-b, where a	AI446563, AI807997, AA872374, AI798452,
			is any integer between 1 to 1357 of	AA256606, AA936249, AI393572, H25408, AW016511,
			SEQ ID NO:1764, b is an integer of	C01415, H28374, AA516090, R43067, AI991488,
			15 to 1371, where both a and b	AA455164, AI217649, AA730296, AI216786,
			correspond to the positions of	AI357214, AI961183, AI537981, AI203429,
			nucleotide residues shown in SEQ ID	AI261590, AI093989, AI950123, R46342, AI803504,
			NO:1764, and where b is greater	AI017015, AA425610, AA535732, AI922416, N21542,
			than or equal to a + 14.	AI805514, R35671, R35782, Z38679, AA258077,
				AI092478, AW170513, AI382468, AA971129,
				AA455366, AA430349, AA090871
1765	HUSGT72	877111	Preferably excluded from the	AA021634, AW028333, AI203234
			present invention are one or more	
			polynucleotides comprising a	
			nucleotide sequence described by	
			the general formula of a-b, where a	
			is any integer between 1 to 752 of	
			SEQ ID NO:1765, b is an integer of	
			15 to 766, where both a and b	

	HPWBM91		a co cue posicions or	
	VBM91			
	WBM91		ide residues snow	
1766 HPW	VBM91		NO:1/65, and where b is greater than or equal to a + 14.	
		877112	1 2	AA496246, AI760599, AI371734, AA476481,
			present invention are one or more	AA496245, AI955212, AI802040, AA628734,
			polynucleotides comprising a	AA476480, AI369165, AI094501, AA744975,
			nucleotide sequence described by	AI609830, AI810354, AI420545, AI381025,
			the general formula of a-b, where a	, AI439413, AI47442
			is any integer between 1 to 722 of	, AA886089,
			SEQ ID NO:1766, b is an integer of	AA505488, AA554685, AA812608, AI125614,
			15 to 736, where both a and b	AA886622, AW389951, AI885739, AA215595,
_			correspond to the positions of	AI000868, AF165185, AF17232
-			nucleotide residues shown in SEQ ID	
			NO:1766, and where b is greater	
			than or equal to a + 14.	
1767 HWI	HWLVB03	877114	Preferably excluded from the	AA112413, AI879634, AI625669, AA287717,
			present invention are one or more	AI027610, AI951403, N51076, AI218397, N72114,
			polynucleotides comprising a	AI924949; AI278323, AI076224, AI921374,
			nucleotide sequence described by	
_			the general formula of a-b, where a	AI221583, AA806202, AI634635, AI357102,
_	•		is any integer between 1 to 507 of	AI272043,
			SEQ ID NO:1767, b is an integer of	AI765676, AW298266, AA768195, AI742632;
_			15 to 521, where both a and b	
			correspond to the positions of	
			$\boldsymbol{\sigma}$	\mathbf{H}
<u></u>			NO:1767, and where b is greater	AA287716, AI424445, N50945, AA127273, H52538,
_			than or equal to a + 14.	AL037272, AA665059, AW340854, AA279150, H10181,
				\$
·				R43464, AW365070, AW079259, Z38935, F03815,
				R40549, AI567606,
	-			AL042450, AL042542, U46349, AL134542, AI433107,
				AL042984, AL043029, U46350, AL043033, AL119497,

				AL119319, AL119483, AL119363,		AL042614, AL119484, AM363220, U46347,
				AW384394,	U46351, Z99396, ALJ	299396, AL134528, AL043011,
				A81671, AR054110,	, 010310, AB026436	11000000 11000000
1768	HAJAM74	877119	Preferably excluded from the	AA026806,	AI243595	
			invention are one c			
			polynucleotides comprising a			
			cribed by			
			wher			
			ween 1 to 439			
			SEQ ID NO:1768, b is an integer of			
			15 to 453, where both a and b			
			correspond to the positions of			
			nucleotide residues shown in SEQ ID			
			NO:1768, and where b is greater			
			equal to a + 14.			
1769	HHMME78	877120	Preferably excluded from the	AA215535,	AA453055, 299396, 1	AL119522, AW392670,
			present invention are one or more	AW384394,	AW372827, AW363220,	, AL119497,
			polynucleotides comprising a	AL119335,	AL119443, AL119319	AL119319, U46349, AL119483,
			nucleotide sequence described by	U46350, A.	U46350, AL119457, AL119324, U46341, AL119484,	U46341, AL119484,
			the general formula of a-b, where a	AL119363,	AL119391, AL036418	AL036418, AL038837,
			is any integer between 1 to 622 of	AL119341,	AL119355, U46351, 1	U46351, AL119496, AL119396,
			SEQ ID NO:1769, b is an integer of	AL037051,	AL036725, AA631969,	, AL036858, U46346,
			15 to 636, where both a and b	AL119418,	AL134524, AL042614	, AL119444, U46347,
			correspond to the positions of	AL134528,	AL042975, AL038509,	, AL039074,
			nucleotide residues shown in SEQ ID	AL119439,	AL037205, U46345, 1	AL134518, AL036924,
			NO:1769, and where b is greater	AL042965,	AL119399, AL134533	, AL042970,
			equal to a + 14.	AI142137,	AL042984, AL119488	, AL042551,
				AL134538,	AL037094, AL037082	, AL037526,
				AL042450,	AL036196, AL037077	, AL037639,
				AL037085,	AL039564, AL042544	, AL043019,
				AL042995,	AL043029, AL134542	, AL042542,
				AL042896,	AL036767, AL036190	, AL043003,
				AL036268,	AL038851, AL038520	, AL038447,

				AW177722, AW378533, AW177728, D58101, D59317,
)5856, DS
				D45273, C14407, AW178781, AI525917, AI557774,
), D59695, C14973,
				AI535686, AW178986, D51221,
				AI525235, C14298, AI
				AW177734, . D80168, AI525242, AW179011, D52291,
				AI525925, AI525912, D51213, AA285331, AI525215,
				C16955, AI525237, D51097, D31458, C05763,
				AI525222,
				T02868, H67858, D80949, C04682, AB028859,
				AJ132110, AR008278, A84916, A62300, A62298,
				AR018138, AF058696, A82595, X68127, AB002449,
				AR060385, X67155, Y17188, D26022, Y12724,
				A25909, A94995, A67220, D89785, A78862, D34614,
				AR008443, I50126, I50132, I50128, I50133,
				D88547, AR066488, AR016514, AR016808, AR060138,
				A45456, A26615, AR052274, X82626, A43190,
				114842, Y09669, A43192, AR038669, AR054175,
				A70867, D50010, AR066490, AR008277, AR008281,
				AR062872, I18367, AR016691, AR016690, U46128,
				I82448, I79511, AR008408, A64136, A68321,
				AB012117, D13509, AR060133, AR066482, AF123263,
				A85396, D88507, A44171, AR032065
1771	HCRNE77	877122	Preferably excluded from the	N46730, N47731, AC005272, AC005826, AC006379,
			present invention are one or more	AC007276, AC004800
			polynucleotides comprising a	
			nucleotide sequence described by	
			the general formula of a-b, where a	
			is any integer between 1 to 720 of	
			SEQ ID NO:1771, b is an integer of	
			15 to 734, where both a and b	
			d to the positions of	
			nucleotide residues shown in SEQ ID	

			NO:1771, and where b is greater		
			than or equal to a + 14.		
1772	HWMBC9	877123	LJ	AA366950	
	4		present invention are one or more		
			polynucleotides comprising a		
			nucleotide sequence described by		
			the general formula of a-b, where a		
			is any integer between 1 to 382 of		
			SEQ ID NO:1772, b is an integer of		
	-		15 to 396, where both a and b		
			correspond to the positions of		
			nucleotide residues shown in SEQ ID		
			NO:1772, and where b is greater		
1773	HWLMS73	877126	Preferably excluded from the	AA527435, AW195324, AI653000, AW051613.	
			present invention are one or more		
			polynucleotides comprising a		
			nucleotide sequence described by	, AW004627, AA397980.	AC002302
			the general formula of a-b, where a		
			is any integer between 1 to 772 of		
			SEQ ID NO:1773, b is an integer of		
			15 to 786, where both a and b		
			correspond to the positions of		
			nucleotide residues shown in SEQ ID		
			NO:1773, and where b is greater		
			than or equal to a + 14.		
1774	HFAMB70	877129		H10992, AL080276	
_			present invention are one or more		
			polynucleotides comprising a		
			nucleotide sequence described by		
			the general formula of a-b, where a		
-			is any integer between 1 to 662 of		
	-		SEQ ID NO:1774, b is an integer of		
			15 to 676, where both a and b		
			correspond to the positions of		
			nucleotide residues shown in SEQ ID		

	W86771											AA595817, H30539, AW022133											AW183176, AI338542, AA687408, AI335604,	, AI741694, AA954272,	AI092736, AI826540, AI675475, AI079357,	AI932722, AW196794, AW028184, AA091428,	AW297724, AI678998				
NO:1774, and where b is greater than or equal to a + 14.	122	present invention are one or more	polynucieotides compilating a	the general formula of a-b, where a	is any integer between 1 to 409 of	SEQ ID NO:1775, b is an integer of	15 to 423, where both a and b	correspond to the positions of	nucleotide residues shown in SEQ ID	NO:1775, and where b is greater	than or equal to a + 14.	Preferably excluded from the	present invention are one or more	polynucleotides comprising a	nucleotide seguence described by	the general formula of a-b, where a	is any integer between 1 to 657 of	SEQ ID NO:1776, b is an integer of	15 to 671, where both a and b	correspond to the positions of	NO:1776, and where b is greater	than or equal to a + 14.	Preferably excluded from the	present invention are one or more	polynucleotides comprising a	nucleotide sequence described by	the general formula of a-b, where a	is any integer between 1 to 1765 of	SEQ ID NO:1777, b is an integer of	15 to 1779, where both a and b	nucleotide residues shown in SEQ ID
	877130											877131											877132								
	HCQAK62											нсорь71											HE9PB28								
	1775											1776											1777								

			NO:1777, and where b is greater.	
1778	HCOCR68	877133	ıΔ	T87566. AW389691. AA505395 P15971 AT.022069
	,		present invention are one or more	14 (114) (1000) (11 (100) (11 (100)) (11 (1000) (11 (1000) (11 (1000) (11 (1000) (11 (1000) (11 (1000) (11 (100)) (11 (1000) (11 (1000) (11 (100)) (11 (1000) (11 (100)) (11 (1000) (11 (100)) (11 (1000) (11 (100)) (11 (10
			_	
			nucleotide sequence described by	
			the general formula of a-b, where a	
			is any integer between 1 to 545 of	
		_	SEQ ID NO:1778, b is an integer of	
			15 to 559, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			-	
			than or equal to a + 14.	
1779	HEPNB10	877134	Preferably excluded from the	AI268381, AI240658, AI302971, W87782, H02333,
			present invention are one or more	315, X828
			polynucleotides comprising a	
			nucleotide sequence described by	
			the general formula of a-b, where a	
			is any integer between 1 to 772 of	
			SEQ ID NO:1779, b is an integer of	
			15 to 786, where both a and b	
-			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
_			NO:1779, and where b is greater	
			than or equal to a + 14.	
1780	HWLNY36	877135	Preferably excluded from the	Z78283, R11554, N44978, AA321699, AA661583,
		_	present invention are one or more	AW275432, AL048969, AI801563, AA640305,
		-	polynucleotides comprising a	AA666295, AA676592, AA483966, AI268826,
•		_	nucleotide sequence described by	AW151247, AW021674, AI174703, AA601376,
			the general formula of a-b, where a	
		_	is any integer between 1 to 674 of	AI370470, H93717, AA846944, C06151, AA469230,
		-	SEQ ID NO:1780, b is an integer of	M77888, AI224583, AI242994, F29968, AA829565,
		-	15 to 688, where both a and b	AA180056, AI090377
		_	correspond to the positions of	AA723132, AA831426, AA525753, AA630476,
			nucleotide residues shown in SEQ ID	AA113757, AA493245, AW275640, AI292275,

	NO:1780, and where b is greater	1, AI457152, T52772, AA233462, AI738
	than or equal to a + 14.	AI309943, AI300597, AW
		AI283329, AA302943, AA720582, AA480486,
		AW087537, AA599069, AI754421, AI474127,
	-	AA601333, AI192465, AA341992, AA367920,
	:	AI583532, AA493789, AW022376, AI053673,
		AA489390, AI417496, T07251, AI797998, AA491743,
_		AA586474, AI590404, D29424, AI538404, AI378950,
		N54538, AI311796, AA084320, AI567676, AI310670,
		AI014332, AA218684, T03928, AL119645, AI282724,
		, N22416, AW264548, AI719298
		, R43468, AA483735, AI349130
		AA666172, AI590442, AI079669, AI654737,
		AA584765, AA228437, AA602105, AI862213,
		AA111897, AI872018, AA847504, AA434165,
		AA342238, AA587835, AI271693, AA368616,
		AW272389, AA347203, AW192199, AA298365,
		AI758981, AL079553, AL078621, AC002055,
		AL096791, AC002316, AL021392, AC005954,
		AC004929, AC000115, AP000518, AC005746,
		AC005011, Z73359, U95742, AC
		Z97632, AL035682,
		AC004587, AL034349, AC007563, Z81450, AC004652,
-		σ
		~
		AC004209, AC004506, AP000514, AL031663,
		AC002554, AC005736, AC002470, AC004834,
		AL035443, AC007564, AC005041, AP000010, Z68273,
		Z97056, AC007308, AF118808, AC004230, AF006501,
		AC004611, AL008716, AL118497, Z84467, Z85986,
		AC005082, AC002310, AC005914, AC005095,
		AC005666, AL078602, AF109907, AC004583,
		, Z82244, AL
		AL034548, AJ003147, AC003685, AC005740,
_		NI.040540 P.0006205 P.0004673 P.0005447

	AC004518, A	AC007110, AL031321, AC004678,
	AL117339, A	AF217403, AC005190, AP000277,
		AC006167, AP000281, AC005251,
	AC003077, A	AP000008, AP000105, AP000037,
	_	Z95113, AP000704, AC002529, AC002465,
		, AP000511
	_	AL049712, AP001053, AL023799,
		1,
		AC020663, AC007066, AC003109, Y15083,
		AC005104, AC006076, AB020859,
	AC007878, A	AC005320, AC004562, AL132799,
	AL023578, A(AC005065, AC006251, AC006275,
	4,	AC004623, AL031223, Z99289, AC006316,
	U91322, AF2	AF207550, AC004477, AC007371, AC006131,
	6	Z99297, Z97832, AL049839, AL133163,
	Z73429, AC0	0
		Z46936, AC005579, AL121767, AC004134,
	AC005015, AI	AP000227, AB004907, U89335, AC005218,
	AC004131, AC	AC006130, AL022322, M94081, AP000087,
		\circ
		AC005924, AC006162, AC004074,
	7.	AJ006997,
		AC004757, AL022725, AC003665,
		3, AC004832
		R
		AC004513, AC004773, AL136295,
		, AC012627
		ý
-		AC005523, AC005261, AC004030,
	_	_
_	AC007225, AI	_
		Æ
		, AC004856,
	_	043, AB009422,
	AP001058, AC	AC005175, AC013256, AC002997,

				AC005594, AC008975, Z68756, L48038, Z75890,
	9704 111111	100	7	TANDANE ABIOAIES
1781	HWLRC68	877137		USSU42, AUZ49/US, AFI84133
			present invention are one or more	
			polynucleotides comprising a	
			nucleotide sequence described by	
			the general formula of a-b, where a	
			is any integer between 1 to 534 of	
			SEQ ID NO:1781, b is an integer of	
			15 to 548, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1781, and where b is greater	
			than or equal to a + 14.	
1782	HWLQM8	877138	Preferably excluded from the	W73224, AI804267, AI379725, AI636783, AI351006,
	′∞		present invention are one or more	H98536, AI365217, N35469, AI219083, AI221578,
			polynucleotides comprising a	AA476333, AI687408, AC007285
			nucleotide sequence described by	
			٦	
			is any integer between 1 to 553 of	
			SEQ ID NO:1782, b is an integer of	
			15 to 567, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1782, and where b is greater	
			than or equal to a + 14.	
1783	HWLMG4	877139	Preferably excluded from the	AI968175, AI970276,
	0		present invention are one or more	
			polynucleotides comprising a	AA516176, AI917709, AI631638, AI625029, AI342081
		_	nucleotide sequence described by	
		_	the general formula of a-b, where a	
			is any integer between 1 to 523 of	
			SEQ ID NO:1783, b is an integer of	٠
			15 to 537, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	

		NO:1783, and where b is greater than or equal to a + 14.	
1784 HWLQ015	15 877140	y excluded from the	, N95228, AI656562, AW
		present invention are one or more	AI375092, AW016802, AI188610, AI985579,
,		the general formula of a-b, where a	
		is any integer between 1 to 600 of	, H19327,
-		SEQ ID NO:1784, b is an integer of	
_	_	15 to 614, where both a and b	, R05523, W69271, Z3
		correspond to the positions of	AA099158, AI984653, AA019723, AI554117,
_		de residue	_
		, and where	AL035700, AC007270
+	\dashv	than or equal to a + 14.	
1785 H2CAC59	9 877142	Preferably excluded from the	AA307078, AA706423, AA994100, AA641669,
		present invention are one or more	AA626714, AA770345, AI360154, AA454000,
		polynucleotides comprising a	AI015598, AI470060, AI470113, AI274091,
		nucleotide sequence described by	AI627230, AI784122, AI563937, AW071839,
		the general formula of a-b, where a	AI937059, AI348119, AI285070, AI401714,
-			4
		SEQ ID NO:1785, b is an integer of	AI139979, AA229891, AI192689, AA745669,
		15 to 495, where both a and b	_
		correspond to the positions of	AI002451, AI568443, AA074240, AA627279,
		ซ	AA451794, R96077, AA767360, AA451795, R96116,
	_		
		than or equal to a + 14.	ò
			9619, D80210, D80240, D80253,
			7, D80212, D80188, D80227, D81030,
			D80391,
	_		D80038, D80022, D80196, D80269, D80164, D59275,
			, AA400769, D80193
-			D50995,
			, AW178893, D81026, D80268
_			F13647, D80949, Z21582, D58253, D80522, D81111,

	אוואפת אפאפבת מאמברנושת פרמובת ברנפרושה
	151 DE2
	MW3/8324, AM314168, AM314167, AW37431, AW177511, AV175816,
-	AW35211
-	D80133, AA285331, AW360811, C1440
	D51097, AW375405, AW360844, D8013
	AW366296, AW360817,
	AW375406, AW378534, AW352171, AW179332,
	AW377672, AW179023, AW377676, AW178905,
	AW179024, D80439, T03116, AW
	AW179020,
-	AW178907, AW179019, AW179018, AW178971, D80247,
	AW352174, D80014, AW179017, AW179004, AW179329,
	AW179012, AW178980, AW17733, AW378528,
	AW178908, T11417, D51103, D80157, AW179009,
	AW178914, AW378543, AW378525, AW367967, T02974,
	D51759, D58246, D58101, AW378539, AW178983,
	AW352120, AW177728, AW178774, AW178781,
	_
	AI557774, T48593, D51213, D4
	AW378533, H67854, AW3679
	A78862, D26022, A25909, D34614, D88547,
	AR025207, X82626, AF058696, AR008278, AB028859,
	\sim
	I19525, A86792, U872
	A94995,
	I50133, I50128, I50126, I50132, AR066488,
	AR060138, AF135125, A45456
	AR066490,
	AR038669, AR066487, I18367, A30438, D88507,
	R054175, D50010, Y17187, AB03
	AR008277, AR008281, A63261, AR064240, AR008408,

				AR062872, A70867, AR016691, AR016690, U46128	, U46128,
				D13509, A64136, A68321, AR060133, I79511, Z32749, U87247, AB023656, AF123263, X9353 APOOR182	9511, X93535,
1786	HWLXJ87	877143	Preferably excluded from the	AW450418, R24589	
			present invention are one or more		
			polynucleotides comprising a		
			the general formula of a-b, where a		
			is any integer between 1 to 570 of		
			SEQ ID NO:1786, b is an integer of		
			15 to 584, where both a and b		
_			correspond to the positions of		
			nucleotide residues shown in SEQ ID		
			NO:1786, and where b is greater		
			than or equal to a + 14.		
1787	HSDSJ26	877145	Preferably excluded from the	AA193531, AI360026, N40228, AA459477,	, N93266,
			present invention are one or more	H85243, AI918187, AI564399	
			polynucleotides comprising a		
-			nucleotide sequence described by		
			is any integer between 1 to 1319 of		
			SEQ ID NO:1787, b is an integer of		
			15 to 1333, where both a and b		
			correspond to the positions of		
			NO:1787, and where b is greater		٠
			than or equal to a + 14.		
1788	HCFBR55	877146	Preferably excluded from the	AI761380,	36, N81076,
			present invention are one or more	AA258570, AA772622,	H22025, AI565200,
			polynucleotides comprising a	AI371499, AA659137, AA879034, AI423953	53,
	-		nucleotide sequence described by	AI084944, U69127	
			the general formula of a-b, where a		
			is any integer between 1 to 536 of		
			SEQ ID NO:1788, b is an integer of		
	_		15 to 550, where both a and b		

correspond to the positions of nucleotide residues shown in SEQ ID NO:1788, and where b is greater than or equal to a + 14.	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 471 of SEQ ID NO:1789, b is an integer of 15 to 485, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1789, and where b is greater than or equal to a + 14.	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 551 of SEQ ID NO:1790, b is an integer of 15 to 565, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1790, and where b is greater than or equal to a + 14.	preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 900 of SEQ ID NO:1791, b is an integer of
correspond to the nucleotide residue NO:1788, and where than or equal to a	Preferably exclude present invention polynucleotides conucleotide sequenc the general formul is any integer bet SEQ ID NO:1789, b 15 to 485, where b correspond to the nucleotide residue NO:1789, and where than or equal to a	Preferably excluded present invention are polynucleotides componies aguence the general formula is any integer betwee SEQ ID NO:1790, b is 15 to 565, where bot correspond to the portion incleotide residues NO:1790, and where the than or equal to a 4	preferably excluded from present invention are or polynucleotides comprision nucleotide sequence describe general formula of a is any integer between 1 SEQ ID NO:1791, b is an
	877147	877148	877149
	HCRNP62	HCRMR04	новне60
	1789	1790	1791

			ond to the position ide residues sind where b		
201	27001		than or equal to a + 14.	- 1	
1/92	HKAUG63	877153	ц.	AA307405, AL037524, AL037501,	AA126654, R97186,
			present invention are one or more	258080	
			eot		
			nucleotide sequence described by		
			eral formula of a-b,		
			is any integer between 1 to 296 of		
			SEQ ID NO:1792, b is an integer of		
			15 to 310, where both a and b		
			correspond to the positions of		
			nucleotide residues shown in SEQ ID		
			NO:1792, and where b is greater		
			than or equal to a + 14.		
1793	H2CBR38	877154	Preferably excluded from the	AA434547, AA278232, AA029146,	AA191433, H00358,
			present invention are one or more	0	
			polynucleotides comprising a	T99622, AA165044, W00839, R35827, AA425497.	827. AA425497.
			nucleotide sequence described by		7. AA449385.
	-		the general formula of a-b, where a	W24857, AA313412, N77971, AW303346, AA455582,	03346, AA455582,
			is any integer between 1 to 1040 of	AI312533, T56653, AA905068, AA304411,	A304411, AW009793,
			SEQ ID NO:1793, b is an integer of	, N77395,	AA129547, AW069049,
			15 to 1054, where both a and b	AI816925, AC002543	
			correspond to the positions of		
			nucleotide residues shown in SEQ ID		
			NO:1793, and where b is greater		
			than or equal to a + 14.		
1794	HRDEW54	877155	Preferably excluded from the	AW303346, AA905068, AW009793,	AA193396,
			present invention are one or more	AA514453, AA587237, AW069049,	AI816925,
			polynucleotides comprising a	AA425497, AA525849, AA455582,	AI309995,
			nucleotide sequence described by	AI768678, AI129597, AA129547,	AI922487, W00839,
			the general formula of a-b, where a	AI679847, AI275507, AW070298,	AI816908,
			nteger between 1 to 783	AA165044,	AA456079,
			O:1794, b is an inte	AI250904, AA405639, AI679273,	AI399923,
			15 to 797, where both a and b	AA600034, AA427915, AA613020,	AA723373,

			correspond to the positions of	AI630755, AA926672, N95773, AI355684, AA576604,
			nucleotide residues shown in SEQ ID	N73000, AI
			NO:1794, and where b is greater	AI125948, AI431758
			than or equal to a + 14.	
				AA449256, AA029146, AA278232, F09333, AA190919,
				H00311, T91257, W02964, N33940, T99623, R49537,
				H8
				AA975401, AW235959, AI767913, Z40018, AA640099,
				AA932232, T49289, T56653, AA029024, T49288,
				W24857,
				AA434547, T49320, AC002543
1795	HBMDC60	877157	Preferably excluded from the	AL031774
		-	present invention are one or more	
	-		polynucleotides comprising a	
			nucleotide sequence described by	
			the general formula of a-b, where a	
			is any integer between 1 to 350 of	
			SEQ ID NO:1795, b is an integer of	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1795, and where b is greater	
			equal to a + 14.	
1796	HOGDM40	877163	Preferably excluded from the	AI459297, AA807285, AA428379, AA443512,
			present invention are one or more	
			polynucleotides comprising a	AI823917, AW296857, R34933, AI964018, R34837,
			nucleotide sequence described by	AL120670, AL120664
			the general formula of a-b, where a	
			is any integer between 1 to 1253 of	
			SEQ ID NO:1796, b is an integer of	
			15 to 1267, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1796, and where b is greater	
			than or equal to a + 14.	
1797	HWLNG61	877165	Preferably excluded from the	

			present invention are one or more	
			\sim	
			nucleotide sequence described by	
			is pur integer between 1 to 440 of	
_				
			15 to 463, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1797, and where b is greater	
			than or equal to a + 14.	
1798	нсост53	877166	Preferably excluded from the	N23022, AI742147, AA399952, AA773713, AI917300,
			present invention are one or more	AA773709, AA768407, N47504, AI339083, AI743525,
			polynucleotides comprising a	AI276208, AI393759, AA933833, H97027, H97002,
			nucleotide sequence described by	AI401278, AI952505, AW294197, AA844082,
			the general formula of a-b, where a	AI990110, AI770034, AI973154, AI381716,
			is any integer between 1 to 877 of	AA620473, AI990671, AA256663, N47503
			SEQ ID NO:1798, b is an integer of	
			15 to 891, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1798, and where b is greater	
			than or equal to a + 14.	
1799	HCRNV59	877167	Preferably excluded from the	AA515852, AA806034, AA642399, AI804718,
			present invention are one or more	AA805516, AI494462, AI478789, AW236212,
			polynucleotides comprising a	AA252353, AI768661, AA721744, AA761615,
			nucleotide sequence described by	AA603497, AL134524, AL134110, AA252268,
			the general formula of a-b, where a	AL047163, AL042898, AL135012, AL042468,
			is any integer between 1 to 420 of	AL042523, AL042420, AL045327, AL045494,
			SEQ ID NO:1799, b is an integer of	AL042741, AL045891, U46344, AL049280, AR066494,
			15 to 434, where both a and b	AL133053, AL122101
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1799, and where b is greater	
			than or equal to a + 14.	
1800	нсорр52	877168	Preferably excluded from the	N94138, AL042183

	W32491, AI557416, AA641955, AC007250	AI432361, AI394416, AI075852, AA479958, AA491075, AA588390, N20112, AW377547, AI888417, AA446881, AF155106, AB033107	AA305314, AI656138
present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 435 of SEQ ID NO:1800, b is an integer of 15 to 449, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1800, and where b is greater than or equal to a + 14.	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 681 of SEQ ID NO:1801, b is an integer of 15 to 695, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1801, and where b is greater than or equal to a + 14.	10 d d 0 d Z d 0 d ×	rat
	877169	877170	877171
	Н FAAH06	HWLMX0 2	HCYBH52
	1801	1802	1803

			present invention are one or more	
•				
		٠	nucleotide sequence described by	
			the general formula of a-b, where a	
			is any integer between 1 to 526 of	
			SEQ ID NO:1803, b is an integer of	-
			15 to 540, where both a and b	
			NO:1803, and where b is greater	
			than or equal to a + 14.	
1804	HCRNX51	877173	Preferably excluded from the	AA232079, AF110400, AB018122
			present invention are one or more	-
			polynucleotides comprising a	
			nucleotide sequence described by	
-			the general formula of a-b, where a	
			is any integer between 1 to 217 of	
			SEQ ID NO:1804, b is an integer of	
			15 to 231, where both a and b	
	_		correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1804, and where b is greater	
			than or equal to a + 14.	
1805	ннерр92	877174	Preferably excluded from the	AI973079, AA813801, AA191593
			present invention are one or more	
			polynucleotides comprising a	
			nucleotide sequence described by	
			the general formula of a-b, where a	
			is any integer between 1 to 374 of	
			SEQ ID NO:1805, b is an integer of	
			15 to 388, where both a and b	
			correspond to the positions of	
-			ro.	
			, and where b	
			than or equal to a + 14.	
1806	HCQAB45	877175	Preferably excluded from the	

ne or more ng a ribed by 1-b, where a 1 to 270 of integer of and b lons of w in SEQ ID greater	the AA305151, H10843 e or more ng a ribed by -b, where a to 320 of integer of and b ons of m in SEQ ID greater	n the AL122007 le or more ing a ribed by l-b, where a ro 907 of integer of and b ons of m in SEQ ID greater	FOCULE OFFICER PRODUCTE CONTRACTOR FOR
present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where is any integer between 1 to 270 ol SEQ ID NO:1806, b is an integer of 15 to 284, where both a and b correspond to the positions of nucleotide residues shown in SEQ NO:1806, and where b is greater than or equal to a + 14.	preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where is any integer between 1 to 320 of SEQ ID NO:1807, b is an integer of 15 to 334, where both a and b correspond to the positions of nucleotide residues shown in SEQ I NO:1807, and where b is greater than or equal to a + 14.	preferably excluded from present invention are or polynucleotides comprision nucleotide sequence describe any integer between 1 SEQ ID NO:1808, b is an 15 to 921, where both a correspond to the position nucleotide residues show NO:1808, and where b is than or equal to a + 14.	Jane Sout
	877176	877181	077101
	HCYBG53	HCQDF43	UCUDIAA
	1807	1808	1800

one or more E17300 ising a escribed by f a-b, where a n 1 to 842 of a and b itions of hown in SEQ ID is greater 14.	rom the one or more ising a escribed by f a-b, where a n 1 to 648 of an integer of a and b itions of hown in SEQ ID is greater 14.	irom the AI769803, AI769743, AI986284, AI031834, alone or more AI017244, AI247689, AI336761, AW445026, alone or more AA933877, AA947886, AI347451, AI344592, alescribed by AI580382, AW302464, AA702771, AA923510, an integer of H66770, H62545, W88899, U66036, AB008164, an integer of AF026303, AJ238392 and b sitions of sitions of is greater 14.	from the AL119483, AL119484, AL119418, AA554958.
present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where is any integer between 1 to 842 of SEQ ID NO:1809, b is an integer of 15 to 856, where both a and b correspond to the positions of nucleotide residues shown in SEQ NO:1809, and where b is greater than or equal to a + 14.	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where is any integer between 1 to 648 of SEQ ID NO:1810, b is an integer of 15 to 662, where both a and b correspond to the positions of nucleotide residues shown in SEQ INO:1810, and where b is greater than or equal to a + 14.	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where is any integer between 1 to 677 of SEQ ID NO:1811, b is an integer of 15 to 691, where both a and b correspond to the positions of nucleotide residues shown in SEQ I NO:1811, and where b is greater than or equal to a + 14.	luded
	877185	877187	877189
	нснѕе50	ноѕруе	HCRMH42
	1810	1811	1812

			polynucleotides comprising a	רפאסטע	AP001053.	AP001053, AF019413, M20903,	, M20903,	
			0	しょうつきつうしな 一	113040011			AC004968,
			nucleotide sequence described by	AC004966				
			the general formula of a-b, where a					
			is any integer between 1 to 601 of					
	_		SEQ ID NO:1812, b is an integer of	-				
	-		15 to 615, where both a and b					
			correspond to the positions of					
			nucleotide residues shown in SEQ ID					
			NO:1812, and where b is greater					
			than or equal to a + 14.					
1813 HS	HSKZE25	877191	Preferably excluded from the	AI740516,	AI739132,	AA631257,	AI741376	ۇ'
			present invention are one or more	AW068935,	AI467852,	AI123717,	AI754551	1,
_	•		polynucleotides comprising a	AI752240,	AW205510,	AA464510,	AW044211,	1,
			nucleotide sequence described by	AW028889,	AW198033,	AI538632,	AA513096	vo
			the general formula of a-b, where a					
			is any integer between 1 to 1191 of					
			SEQ ID NO:1813, b is an integer of					
			15 to 1205, where both a and b					
	•	_	correspond to the positions of					
			nucleotide residues shown in SEQ ID					
			NO:1813, and where b is greater					
			than or equal to a + 14.					
1814 HC	HCRMP38	877194	Preferably excluded from the	AI623320,	AL023654			
			present invention are one or more	-				
			polynucleotides comprising a					
			nucleotide sequence described by					
			the general formula of a-b, where a					
			is any integer between 1 to 586 of					
			SEQ ID NO:1814, b is an integer of					
			15 to 600, where both a and b					
			correspond to the positions of					
			nucleotide residues shown in SEQ ID					
			NO:1814, and where b is greater					
			than or equal to a + 14.					
1815 HE	HDPXD55	877195	Preferably excluded from the	AL110186,	AB011097			

			1		
			polynucleotides comprising a		
			nucleotide sequence described by	-	
			the general formula of a-b, where a		
			is any integer between 1 to 551 of		
			SEQ ID NO:1815, b is an integer of		
			15 to 565, where both a and b		
			correspond to the positions of		
			nucleotide residues shown in SEQ ID		
			NO:1815, and where b is greater		
			than or equal to a + 14.		
1816	HHMMB4	877200	Preferably excluded from the		
	0		present invention are one or more		
			polynucleotides comprising a		
			nucleotide sequence described by		
			the general formula of a-b, where a		
			is any integer between 1 to 272 of		
			SEQ ID NO:1816, b is an integer of		
			15 to 286, where both a and b		
			correspond to the positions of		
			de residues :		
			NO:1816, and where b is greater		
			than or equal to a + 14.	:	
1817	HEQAN41	877202	Preferably excluded from the	AW003740, W81689, AI862673, P	AW270849, AI912038,
			present invention are one or more	AI703038, AA937086, AI279103,	AA282925,
			polynucleotides comprising a	AI078559, AI768831, AA313607,	AI275886,
			nucleotide sequence described by	AI432429, AA903131, AI870642,	AI189825,
		_	the general formula of a-b, where a	AA283134, W81688, AI521151, A	AW044071, AA410488,
			is any integer between 1 to 1306 of	AA827169, AA730751, AA256352,	AW131390,
			SEQ ID NO:1817, b is an integer of	AI970675, AA989435, AA918065,	AI813309,
			15 to 1320, where both a and b	AI969627, AA255498, AA621557,	AA828340,
			correspond to the positions of	AI693110, AI351613, AI471645,	AA025513,
			nucleotide residues shown in SEQ ID	AI912910, AA410307, AW071626,	AI655122,
			NO:1817, and where b is greater	AI800296, AI651526, AI368793,	AA976771,
			than or equal to a + 14.	AI631084, AI829747, AI620149,	AI970920,
				AA256209, AI422613, AI826838,	AW389929,

CHORTH NOSCOREN OFFICER	6, AAS62664, AL91/US AT859906 AT915081	AI279417, AA678616,	7, AA832016, AI922614,	2, AA484892, AA610255,	AA609826, AI631059, AI797998, AI869786, F08655,	AA598605, AI038324, AA857812, AI018726,	AA807579, AA778962, AW265688, AW019964,	AA904211, AI383596, H59611, AI150934, H59651,	, AW078821, AW390284	~	AI472736, F33820, AW440568, R99613, AA678932,), T76991,	AW270351, AA362791, AI803741, AI889995,	AI359200, AA126814, AI419337, AI361090,	AA345594, AW192518, AI671077, AW026305,	, H39839, AW	AA643829, AA402113, AI289050, AA653291,	AA436140, AI358776, F17537, AI284092, H38901,	AI291419, AA484022, AF003627, AF035397,	AF086459, AF130357, AC007656, AF111169,	AC005231, AC002316, AP000350, AP000045,	AL049830, AC004820, AL133448, AC004990, Z49258,	AC007055, AL121603, AL031984, AC006084, L78810,	Z82208, X51956, AL031602, U47924, U85195,	AC003029, AE000658, AC006251, AC005696,	AC007878, AL049692, AC005480, AC005082,	AC000379, AC007057, AL049872, AC005006,	AL031433, AC005484, AL031295, AC007687,	. AC005089, AL096791, AC002312, AL050305,	AC006443, AL031728, AL133371, AC002432,	5, AC00533(AC002365, Y10196, AC004408, AC005212, AL022240,
																-			 -													

AL049694, AC005048, AC005902, AC010205,
AC004383, AL049553, AC004148, AF064866,
AC003982, AF196779, AL049641, AC008041, L44140
AF095901, AL050404, AL031293, AF207550,
AJ003147, AC005778, AC003101, AC005695,
AL121652, AC006359, AL024498, AP000113,
AC003107, AP000352, AC000026, AC004675,
Ę
, AC005209, AC003036,
3, AP000247
AC005971, Z95115, AL034377, AC
Z69715, AP000304, AL
AC000031, AF053356,
l, AC007350, AC005102,
D84394, AC005943, AC003973, AC004685, AC007014
AL035405
AC004447, AC004815,
AC005015, AC007686, AC004638, Z73988, AC004230,
C004883, AC
AL050347, AC009330,
, AC008372, AC005726,
5, AC003692, AL035697,
6, AL031774, AL035455,
, AL022719, AC002115
, AC004477, A51133, A7
AC005907, AC005519, AL121782, Z98742, AP000030,
AC005365, AL008729, AF217403, AL132985,
AC005562, AC004890, AC006948, AC002551,
AC004185, AC005844, AL035403, AC004539,
AP000115, AP000695, AC009247, AL031730,
AC002429, AL109963, AL033523, AC000112,
AC007263, AL133245, AL031053, AL021397,
בניטניסטא סטטונטוא טנטנימא נדטנטטטא

				AL122020, AL021154, AC005666, AL136295,
				AL009179, AL022721,
				AC005874,
				AC005969, AC006160, AL133244, AC002550,
				AL022313, AI632057
1818	HSDZB30	877205	Preferably excluded from the	AA129439, AA425398, AI381416, R17127, AI418660,
			present invention are one or more	AA314750, F32787, AI590092, AW021547, AA151302,
			polynucleotides comprising a	Z42142, AA904204, U77327, AF064105
			nucleotide sequence described by	
			the general formula of a-b, where a	
			is any integer between 1 to 807 of	
			SEQ ID NO:1818, b is an integer of	
-			15 to 821, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1818, and where b is greater	
			Ψ	
1819	HWLWHS	877206	Preferably excluded from the	AI989601, AC005593
	9		present invention are one or more	
			polynucleotides comprising a	
			nucleotide sequence described by	
			the general formula of a-b, where a	
			is any integer between 1 to 356 of	
			SEQ ID NO:1819, b is an integer of	
			15 to 370, where both a and b	
			nd to the positions of	
			nucleotide residues shown in SEQ ID	
			than or equal to a + 14.	
1820	HWLOT46	877207	Preferably excluded from the	
			present invention are one or more	
			polynucleotides comprising a	
			nucleotide sequence described by	
			the general formula of a-b, where a	
			is any integer between 1 to 388 of	

			SEO ID NO:1820, b is an integer of	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1820, and where b is greater	
			than or equal to a + 14.	
1821	HOVCR67	877208	Preferably excluded from the	
			present invention are one or more	
			polynucleotides comprising a	
			nucleotide sequence described by	
			the general formula of a-b, where a	
			is any integer between 1 to 334 of	
			15 to 348, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1821, and where b is greater	
			than or equal to a + 14.	
1822	HLHSV54	877211	Preferably excluded from the	
			present invention are one or more	
			polynucleotides comprising a	
			nucleotide sequence described by	
			the general formula of a-b, where a	
			is any integer between 1 to 498 of	
	_		SEQ ID NO:1822, b is an integer of	
			15 to 512, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
_			NO:1822, and where b is greater	
			than or equal to a + 14.	
1823	HSYBZ84	877212	Preferably excluded from the	AA922141, AA505358, AA515537, AI439152,
			present invention are one or more	AA603688, AI279253, AI003069, H09774, R61798,
			polynucleotides comprising a	N46444, N48945, R45147, Z45425, R55783, R43907,
			nucleotide sequence described by	R14995, AA348815, AB032971
			ral formula of a-b,	
			is any integer between 1 to 926 of	

	·		SEQ ID NO:1823, b is an integer of 15 to 940, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1823, and where b is greater than or equal to a + 14.	
1824	H2LAC34	877213	10 d d 0 d 2 0 0 d ·	AA304651, AI372785, AA496464, R09787, D59627, C16955, D45273, D80168, D52291, D51213, T03048, D59695, C14298, D51079, D80949, D80258, Z33452, AW360780, D59503, C14407, D58246, D80014, C14227, D80064, AI535686, D81111, T11417, T02974, AW377669, D58101, D52059, H67854, D59317, D80038, H67866, AI525216, AI52528, AA809122, AA305578, D50979, D80195, D52317, C15076, D80193, D80251, D59551, C06015, D81026, D80269, D80022, D59467, D80164, D59275, D80045, D80269, D80022, D59467, D80164, D59275, D80045, D80269, D80022, D59467, D80164, D59275, D80045, D80269, D80210, D80269, D80210, D80269, D80247, D51030, D80188, D57483, C3092, D80243, D80157, D51103, D59859, C14331, D80212, D80268, D80247, D51022, D80196, D59619, D80133, D80247, D51022, D80196, D59619, D80133, D80247, D51022, D80196, D5927, D31458, D80248, D59610, C14014, D51221, Z30160, D80522, AA514184, AI525903, AI525923, AI525215, T11191, AI525237, AI525903, AI525912, AI525215, T11191, AI525237, AI525917, AR51580, AB010386, AR060385, AA132110, AB028859, AB010386, AR060385, AA132110, AB028859, AB010346, AR060385, AH1442, I79511, AR008278, U37689, I81198, AR01889, AB010348, AR008278, AR008377, AR008281

				APOTRISA	AFASAGA	D47134	
1825	HCOAE29	877214	Preferably excluded from the		AA730263		
	1771177	4			50305		
			present invention are one or more				
			polynucleotides comprising a				
			nucleotide sequence described by				
			the general formula of a-b, where a				
			is any integer between 1 to 627 of				
			SEQ ID NO:1825, b is an integer of				
			15 to 641, where both a and b				
			correspond to the positions of				
		_	nucleotide residues shown in SEQ ID				
-			NO:1825, and where b is greater				
			than or equal to a + 14.				
1826	HCRMV19	877215	Preferably excluded from the	N72981			
			present invention are one or more				
-			polynucleotides comprising a				
			nucleotide sequence described by				
			the general formula of a-b, where a				
-			is any integer between 1 to 433 of				
			SEQ ID NO:1826, b is an integer of				
			15 to 447, where both a and b	,			
			correspond to the positions of	•			
			nucleotide residues shown in SEQ ID				
			NO:1826, and where b is greater				
			than or equal to a + 14.		•		
1827	HWLMF31	877218	Preferably excluded from the	AI806805,	AA909734,	AI205805,	AI208930,
			present invention are one or more	AI023837,	AI024558,	AA808303,	AI239842,
			polynucleotides comprising a	AA904642,	AI200741,	AA861427,	AI808962,
			nucleotide sequence described by	AA971918,	AA806642,	AC004542	
			the general formula of a-b, where a				
			is any integer between 1 to 576 of				
			SEQ ID NO:1827, b is an integer of				
			15 to 590, where both a and b				
			correspond to the positions of				
			de residues show				
			NO:1827, and where b is greater				

			than or equal to a + 14.	
1828	HFIIZ28	877220	erably ent inv nucleot eotide general ny inte ID NO:1 o 425, espond eotide or equ	88, AI418599, AI151240, AI80 48, AA878931, AI241082, AA93 73, AA194942, N30395, AA5237 68, AI472706, AI336385, AI28 97, AI754786, AW085594, AA87 50, AL044439, AA180129, AA52 83, AA628042, AA627935, AA91 91, AI289442, AL034430
1829	нсорк28	877222	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 368 of SEQ ID NO:1829, b is an integer of 15 to 382, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1829, and where b is greater than or equal to a + 14.	N75183, AI366031, F12542, T74151, AC012627
1830	ннео129	877229	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 818 of SEQ ID NO:1830, b is an integer of 15 to 832, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1830, and where b is greater	AA446316, AA446497, AI198963, H38387, AI444827

	than or equal to a ± 14	
Τ,		
057/0	ο.	
_	present invention are one or more	AW130854, AI092715,
	5	AA846295, AI027808
-	nucleotide sequence described by	N33620, AI215790, AI
—–	al formula of a-b,	AA771890,
	reger between 1 to 576	
	SEQ ID NO:1831, b is an integer of	AA040794,
	correspond to the maritimes of	
nu	nucleotide residues shown in smorth	
8	and where b is greater	
-	equal to a + 14.	
877231 Pr	Preferably excluded from the	AL135440, W20119, AI810591, AT089310 AB044704
pre	present invention are one or more	3. AA039903. AT420778
od	polynucleotides comprising a	, AI093762, AI982907.
nn.	nucleotide sequence described by	AI246659,
the -	e general formula of a-b, where a	AI088688,
18	is any integer between 1 to 3252 of	AI023926,
SES	SEQ ID NO:1832, b is an integer of	, AA669926, AA523605,
7	Is to 3266, where both a and b	AI569996, AI354883,
COL	d to the positions of	AI168582, N3
חמכו	nucleofide residues shown in SEQ ID	AI
	, and where b	AI636575, AA214649, W81054
	chan or equal to a + 14.	AA723161, R70656, AI086670,
_		AW262560, W02383, AA906264,
		AI276236, AI141343, AA868115
		AA862839, AI275375, H10905, AA129975, R80462,
		A846612, AA
		AI309745, AA359784
		, AI990659, AA379173, Z40721,
		, R60952, AA670197, AA435840,
		~
		AI206251, AI476295, AA211075, AI619485, N90439,

				R05760. A	AA079305. WC	W07456. AA0	AA079306. AA84	AA847920.
				AW387693,		AI689470,	A1953765,	
				AI470293,	AA806719,	AA631120,	AI889818,	
				AI274527,	AI249962,		AI888621,	
				AI365256,	AI679095,	AW149876,	AF003626,	Y10043,
				AF022465,	Z83826, Z	93931, ACO(Z83826, Z93931, AC002526, Y10044	144,
				AC005479,	AL024505,	AL034450,		
				AL049709,		AF047701,	L05085, AC004493	3004493,
				AF026008,	Z20724, Z	Z20735		
1833	HWMBOS	877232	Preferably excluded from the	AI289115,	AA653396,	AI280875,	AW439596,	
	0		present invention are one or more	AA147044,	AI683907,	AI186619,	AW191991,	
			polynucleotides comprising a	AI422310,	AI653662,	AA825197,	AA854077,	
			nucleotide sequence described by	AA916637,	AA810755,	AI624228,	AI763289,	AA449797
			the general formula of a-b, where a					
-			is any integer between 1 to 844 of	<u>-</u> -				
			SEQ ID NO:1833, b is an integer of					
			15 to 858, where both a and b					
			correspond to the positions of					
			nucleotide residues shown in SEQ ID					
			NO:1833, and where b is greater					
			than or equal to a + 14.					
1834	нсовр64	877233	Preferably excluded from the	AW008122,	AC005021,	L48431		
			present invention are one or more					
			polynucleotides comprising a					
		_	nucleotide sequence described by					
			the general formula of a-b, where a					
			is any integer between 1 to 283 of					
		_	SEQ ID NO:1834, b is an integer of					
			15 to 297, where both a and b					
			correspond to the positions of					
			nucleotide residues shown in SEQ ID					
			NO:1834, and where b is greater					
			than or equal to a + 14.					
1835	HATAP30	877234	Preferably excluded from the	AI828084,	AW292950,	AI955290,	AI425012,	D54798,
			present invention are one or more	AA101714,	AA661732,	AI082095,	AI433898,	N78571,
			polynucleotides comprising a	AA563807,	AI457762,	AA460668,	AA101715,	

MI148116, AI276830, AI378227, AI148121, AI082653, AI972872, AA631712, AI272196, 4 of AA603075, AI018047, AI453834, AI223254, Of AI026628, AW298807, AI280067, AI378917, T19338, T33356, AA761507, AI272883, R51104, AA644592, T03688, AI274939, AI268664, AI690246, T33873, Q ID N52587, AA461016, T32236, AA464590, AA693417, AA18879, T15930, AL120494, AA371748, N75010, R41316, R41317, AI834293, D81373, AA767242, AW386979, R42324, T33358, T3357, AI366186, T27271, W01584, AI700577, AI767391, AI760808, W26393, W07166, AA861382, AI816326, AI291384, AI913952, W05753, AA488932, AA411945, T09288, R11766, H24112, AW293062, AI277039, R18459, U66702, U81561, U65065, U73458, A633346, A63355, AF007555, Y08569, A63357, U91574, U82439,	AA316077, AW407693, R35424, AL121134, AA356852 re F12867, AA776842, AW163365, M74089 y of of Of	AA402106, AI734033, AA401995, AI821646, AW438 re
nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 1244 of SEQ ID NO:1835, b is an integer of 15 to 1258, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1835, and where b is greater than or equal to a + 14.	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where is any integer between 1 to 747 of SEQ ID NO:1836, b is an integer of 15 to 761, where both a and b correspond to the positions of nucleotide residues shown in SEQ INO:1836, and where b is greater than or equal to a + 14.	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by
	877235	877237
	H2LBB51	H6EDT19
	1836	1837

	W53026, AF180919	W92133, AL035400	AA307110, AI791261, N36579, D80195, D59467, D80164, C15076, D80227, D80269, D59275, D59502, D58283, D59859, D80022, C14331, D80166, D51799, D51423, D59619, D59610, D80210, D80391, D80240, D80253, D80043, D59787, D81030, D80038,
is any integer between 1 to 911 of SEQ ID NO:1837, b is an integer of 15 to 925, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1837, and where b is greater than or equal to a + 14.	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 528 of SEQ ID NO:1838, b is an integer of 15 to 542, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1838, and where b is greater than or equal to a + 14.	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 428 of SEQ ID NO:1839, b is an integer of 15 to 442, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1839, and where b is greater than or equal to a + 14.	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a
	877240	877242	877247
	HWLOW8	HWLMB22	H2CBA14
	1838	1839	1840

	is any integer between 1 to 501 of	AA305409, D80378, D80212, D80366, D50979,
	NO:1840, b is an	D80193, D80196, D80188, D80219, D59927, D57483,
	15 to 515, where both a and b	5, D59889, D80241, C14389,
	correspond to the positions of), C75259,
	nucleotide residues shown in SEQ ID	AW378532, AW178775, AI732942, AA305578,
	NO:1840, and where b is greater	AW179328, D80134, AW177440, D81026, D51250,
	than or equal to a + 14.	8
		D80522, F13647, D80268, AW378540, D80168,
_	-	, C14298
		C14227,
		Z21582, AW377671, D81111, AW360834, AA514186,
		, AW375405, D80132
		D80247, AW360817, AW375406, AW178905, AW378534,
		_
		AA285331, AW179020, AI557751
		AW177456, C06015, D51097, AW352170, AW177731,
		D51103, AW179019, AW179018, T03116, D80157,
		AW378528, AW178908, AI557774, AW352174,
		AW352163, D80258, AI525923, D80014, T48593,
		D59627, AW178774, AW378539, D45260, AA809122,
		T11417, H67866, D45273, C03092, H67854,
		AW367950, AI525227, D51213, AW178986, D59317,
		D59503, T02974, D58246, C14973, AI525917,
		1221, D59474, AI525920, D59551,
		1, Z30160, H67858, AI525925, AI52523
		AI525242, T02868, Z33452, AI525239, C16955,
		AI525912, AI525237, AI525215, AW378542, C13958,
•		D31458, A84916, AJ132110, A62300, A62298,
		AR018138, X67155, Y17188, D26022, A25909,
		D88547, AR008278, AB028859, X82626, AR025207,
		A82595, 'Y12724, A94995, AR060385, AB002449,
		AB012117, AR066482, X68127, AR008443, A85396,

				IS0126, IS0132, IS0128, IS0133, A44171, A85477,
	-			U87250, AR066488
				X93549, AR060138, A45456, A26615, AR052274,
				I14842, Y09669, A43192, A43190, AR038669,
		_		AR066487, AR054175, A30438, Y17187, I79511,
				I18367, A63261, D50010, AR008277, AR008281,
				U46128, AR008408, AF135125, A64136, A68321,
				D13509, AR060133, U87247, AB033111, AR064240
1841	HCRNM80	877250	Preferably excluded from the	AI479603, AW190581, AA573923, AA883422,
			present invention are one or more	AI031618,
			polynucleotides comprising a	AI332605, AI738984, AA910770, N30717, AA146619,
			nucleotide sequence described by	AI348584, AA309589, AA143550, AA146653,
	-		the general formula of a-b, where a	AW293078, AA625575, AA625979, AA676991,
			is any integer between 1 to 1013 of	AW384713, AA494197, AA679394, AA085095,
	-		SEO ID NO:1841, b is an integer of	AI800002, AI739098, AI126129, N41331, AI682193,
	•			A143647, H79815, AA62
	•		-5	AI372964, C05152, N75441, AA085143, W89067,
			nucleotide residues shown in SEQ ID	AI290775, AI202571, T99951, AW008713, W95658,
			NO:1841, and where b is greater	AW384743, R45400, AI201781, AW389792, AW389779,
			than or equal to a + 14.	AW389790, W95657, AA721631, AA354111, AW389774,
			•	R29667. AW389836.
				AI267185
1842	HCOCC04	877251	Preferably excluded from the	N65940, H82959, H72780, R09098, H90731
			present invention are one or more	
-			polynucleotides comprising a	
			nucleotide sequence described by	
			the general formula of a-b, where a	
			is any integer between 1 to 430 of	
			SEQ ID NO:1842, b is an integer of	
			15 to 444, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1842, and where b is greater	
			than or equal to a + 14.	
1843	HCQCI17	877254	Preferably excluded from the	AA129983, M73489, S57551, D17513, Z74734

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			present invention are one or more polynucleotides comprising a	
			nucleotide sequence described by	
			the general formula of a-b, where a	
			is any integer between 1 to 536 of	
			SEQ ID NO:1843, b is an integer of	
			15 to 550, where both a and b	
			correspond to the positions of	
			۲.	
			NO:1843, and where b is greater	
			than or equal to a + 14.	
1844	HFIYJ63	877255	Preferably excluded from the	AL135394, W87908, AB002331
			present invention are one or more	
			polynucleotides comprising a	
		_	nucleotide sequence described by	
			the general formula of a-b, where a	
			is any integer between 1 to 312 of	
			SEQ ID NO:1844, b is an integer of	
	•		15 to 326, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1844, and where b is greater	
			than or equal to a + 14.	
1845	HWLOW5	877256	Preferably excluded from the	H23330, AI796906
	_		present invention are one or more	
			polynucleotides comprising a	
			nucleotide sequence described by	
			the general formula of a-b, where a	
			is any integer between 1 to 563 of	
-			SEQ ID NO:1845, b is an integer of	
			15 to 577, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1845, and where b is greater	
			than or equal to a + 14,	
1846	HHFBA07	877257	Preferably excluded from the	AW130559, AA604942, AI125644, AI703464,

			present invention are one or more	AW103052, AI391708, AI452537, AI460380,
	-		polynucleotides comprising a	AI050784, AI949725, AI052071, AW237646,
			nucleotide sequence described by	AI538701, AI435508, AA621302, AA233121,
			the general formula of a-b, where a	AI348838, AI339780, AI800246, T67212, AI144461,
			is any integer between 1 to 718 of	AW130699, AA527371, AW205441, AA346401,
			SEQ ID NO:1846, b is an integer of	AI247525, AI352551, AI651506, AA707110, R46530,
			15 to 732, where both a and b	AI927033, AI560516, R46529, AI918364, N75541,
			correspond to the positions of	R51933, R72231, H45846, T67213, AA627945,
			nucleotide residues shown in SEQ ID	N40063, AA233205
			NO:1846, and where b is greater	
			than or equal to a + 14.	
1847	HWLD051	877258	Preferably excluded from the	AI830540, AA357636, AAS16122, AI391596,
			present invention are one or more	AI670727, AA814145, AA661893, AA554670,
			polynucleotides comprising a	AI335153, AW157547, AI862260, D31492, AA992253,
			nucleotide sequence described by	AA972187, AI271839, AI218276, AC005606, AC005363
			the general formula of a-b, where a	
			is any integer between 1 to 302 of	
			SEQ ID NO:1847, b is an integer of	
			15 to 316, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1847, and where b is greater	
			than or equal to a + 14.	
1848	HLSAE05	877261	Preferably excluded from the	AA307126, Z99396, AW392670, AW372827, AW384394,
			present invention are one or more	
			polynucleotides comprising a	AL119363,
			nucleotide sequence described by	AL119457, AL119324, AL119483, AL119484,
			the general formula of a-b, where a	AL119391, AL119341, AL119355, U46350, U46349,
			is any integer between 1 to 703 of	AL119396, U46351, AL119418, AL036418, AL038837,
			SEQ ID NO:1848, b is an integer of	AL037051, AL036725, AA631969, U46346, AL119444,
			15 to 717, where both a and b	U46347, AL042614, AL042965, U46345, AL134518,
			correspond to the positions of	AL036858, AL134533, AL042970, AL134524,
			nucleotide residues shown in SEQ ID	AL119439, AL037205, AL134528, AL042975,
		_	NO:1848, and where b is greater	AL119401, AI142137, AL119399, AL036924,
			than or equal to a + 14.	AL042984, AL042551, AL134538, AL042433,
				AL042995, AL119320, AL042850, AL119488,

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				2102010	0 2 7 7 7 7 8	010110	000000
				AL036303,	ALO42430,	AL045019,	AD043023,
				ALU3/085,	ALU42544,	AL042542,	ALU42896,
				AL037094,	AL037526,	AL036196,	AL037639,
				AL119304,	AL043003,	AL036268,	AL037082,
				AL036767,	AL037077,	AL036190,	AL119464,
				AL036774,	AL038520,	AL036998,	AL038851,
				AL038447,	AL036733,	AL037178,	AL036238,
-				AL036719,	AL037615,	AL037027,	AL036765,
				AL036191,	AL036679,	A81671, A	45
				AR023813,	AR064707,	AR069079,	AR054110, AB02643
1849	HCRPJ05	877263	Preferably excluded from the				
			present invention are one or more				
			polynucleotides comprising a				
			nucleotide sequence described by				
			the general formula of a-b, where a				
			is any integer between 1 to 349 of				
			SEQ ID NO:1849, b is an integer of				
			15 to 363, where both a and b				
			correspond to the positions of				
			nucleotide residues shown in SEQ ID				
			NO:1849, and where b is greater				
			than or equal to a + 14.				
1850	HCYBD05	877264	Preferably excluded from the	AA305049,	N50596, A	AL120893, US	U55937, U81001
			present invention are one or more				
			polynucleotides comprising a				
			nucleotide sequence described by				
			the general formula of a-b, where a				
			is any integer between 1 to 522 of				
			SEQ ID NO:1850, b is an integer of				
			15 to 536, where both a and b				
			correspond to the positions of				
			nucleotide residues shown in SEQ ID				
			NO:1850, and where b is greater				
			than or equal to a + 14.			•	
1881	HKLSD44	877272	Preferably excluded from the	AI183955,	AW136574,	AI654355,	D13902, D13897,
			present invention are one or more	L25648, A	AC007993, D.	D13899, M17523	523, S57220,

			polynucleotides comprising a	L37369, Z58904
			al formula of a-b, when	
			teger between 1 to 522	
			SEQ ID NO:1851, b is an integer of	
			15 to 536, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1851, and where b is greater	
			than or equal to a + 14.	
1852	HFIXP45	877274	Preferably excluded from the	U69202, AI341555, AI808490, AI347923, AA903736,
			present invention are one or more	AA210763, AI139380, AI631374, AA129554, W70085,
			polynucleotides comprising a	AI648656, AA932877, AA136568, R39447, F09386,
			nucleotide sequence described by	AI351322, AW001825, T77200, F11728, T09089,
			the general formula of a-b, where a	T10129, H17528, T10128, AI867156, R59448,
			is any integer between 1 to 1991 of	R59388, AI868687, Z19406, AI474036, Z42465,
			SEQ ID NO:1852, b is an integer of	Z28503, Z38662, F06906, F04874, R13169, H17840,
			15 to 2005, where both a and b	AA348361, R13170, Z45682, AB000814, D89722,
			correspond to the positions of	U60415, AF044288, AB000812, AB000813, AB012600,
			nucleotide residues shown in SEQ ID	U51627, AF015953, AB012601, AB015203, AB012602,
			NO:1852, and where b is greater	AB014494, AF070917, AB000815, AB000816
			than or equal to a + 14.	
1853	HAQNS64	877275	Preferably excluded from the	AC005740
			present invention are one or more	
			polynucleotides comprising a	
			nucleotide sequence described by	
			the general formula of a-b, where a	
			is any integer between 1 to 552 of	
			SEQ ID NO:1853, b is an integer of	
			15 to 566, where both a and b	
			correspond to the positions of	
			NO:1853, and where b is greater	
			than or equal to a + 14.	
1854	нсордо	877280	Preferably excluded from the	AW404075, AA469906
			present invention are one or more	AA316159, N42495, R57922, Z59290

			polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 236 of SEQ ID NO:1854, b is an integer of 15 to 250, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1854, and where b is greater than or equal to a + 14.				
1855	нсдсрв1	877281	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 1145 of SEQ ID NO:1855, b is an integer of 15 to 1159, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1855, and where b is greater than or equal to a + 14.	AI207647, AI065011, AI207442, AI207442, AI207715, AL047029, AL047029, AL047022, AMO82028, AI051094, AI051094, AI051094, AI051094, AI051094, AA555071, AA55607, AA55607, AA558890, AA10807, AA558834, AA224754, AA224754, AA897022, AA109069, AA109069,	AIO65109, AII33300, AII14870, AII13620, AII10641, AA401001, AA53278, AA516319, AA554486, AII74849, AA554486, AII74849, AA564569, AA16123388, AA583220, AA583220, AA192604, AA583220, AA583220, AA583220, AA583220, AA583220, AA583220, AA583220,	AI207735, AI13323 AI110723, AI13291 AI064757, AI13291 AI13496, AA29304 AA149787, AI82743 AA149787, AIR2743 AA196910, AA554914, AA563936, AIS57100 C18953, AA654914, AA196910, AA55411, AA196910, AA55411, AA595757, AA14967 AA595757, AA14967 AA595757, AA14967 AA595757, AA14967 AA595503, AA54833 AA179156, AI13316 AA595503, AA512990 AA524960, AM36863	AI133231, AI132917, AI132917, AI132917, AI132979, AI132979, AI132979, AI132970, AI149240, CI7649, AA663700, AI557108, AA554914, AA534001, AA554113, AA554113, AA554113, AA564914, AA53401, AA502430, AA502430, AA512996, AA886497, AA886497, AA886497, AA886497, AA886497, AA886497, AA886497, AA893439, AA593439, AA593439, AA593439, AA566006, AA566006,
				AA579806, AA587814,	AA235499, AI535677,	AA576180, AW368637,	AA834302, AA400809,

			758,											930,					•		-			290,					712,			966			
AA653974,	AA464752,	AI025574,	AI910010, AA508758	AA481923,	386075,	AA526743,	AA985612,	AA431814,	AA709167,	AI783446,	AA583092,	AW070565,	AA091624,	AA886562, C039	AI015676,	AA992091,	AI680484,	AA079806,	AA568749,	R28950, C18721,	AA649597,	AA632764,	AA181000,	C15091, AW382!	AA194421,	AW364429,	AW373685,	AA554414,	AI862143, AI908712	AW178905,	AA714432,	AI935127, X628	038112,	AF014888,	AF014897
AA617685, AA6	AA428850, AA4	-	AA143135, AI910	AW373400, AA4	AA554801, AA88	AA879019, AAS	AA090685, AAS	AA564658, AA4	AI954125, AA7	AA532797, AI7	AI910011, AAS	AA680242, AW	AA620694, AAC	AI133009, AA8	AA194368, AIO	AI936914, AAS	AA694521, AI6	AW371871, AAC	AI620133, AA	C03144, R2895	AA095478, AA6	AW178904, AA6	AA916453, AAJ	_	_	_	AI253336, AW3	ο,	H01671, AI862	AW367539, AW1		AA090224, AIS	, J01415, I	AF014883, AFC	AF014892, AFC
AA593495, AA	AA725126, AA	AA291811, AA	C18039, AA14	_	AA923266, AA	AA938043, AA	AA554076, AA	AA112939, AA	_	AA086336, AA	AA470370, AJ	AW371295, AA	AI910004, AA	AA453608, AJ		AA877931, AI	_	AA934835, AW	AA724218, AJ	AA456614, CC	AA506494, AA	AA630561, AV	AA196736, AA	_	_	_	AW373663, Al	AW364463, AV	AI004318, HC	AI565446, AM		AI833081, A	o		AF014890, AF
AA079632, 1	AA523492, 1	AA507391, 1	AA834333, (AA527764, 1	AA582805, 1	AA908596, 1	AW378088, 1	AA595582, 1	AA401126, 1	AA171612, #	AA576154, 1	AA564029, 7		AA086135, A	AA464751, A	AA176484, 1	AA708229, 1	AW175960, 1	AA650245, 1	AI525240, A		AA534145, 1	_	AA127860, A	_	7,	AW373695, 1	AI832579, A	AA159642, A	AI052019, A	AA193076, 1		X93334, M1(AF134583, A	AF014889, A
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				AF014898, AF014901, AF014893, AF014894, AF014899, AF014891, AF014895, D38116, D38113, X93335, AF014903, AF014904, AF014917, AF014910, AF014920, AF014908, AF014913, X93347, AF014905, AF014916, AF014906, AF014907, AF014909, D38114, AF014902, AF014919, X97707, D38115, D38484, X99256, X89843, U95646, X14848, X59268, S75895
1856	HLHE146	877282	preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 922 of SEQ ID NO:1856, b is an integer of 15 to 936, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1856, and where b is greater than or equal to a + 14.	AI669644, AI925693, AA548892, AA233718, AI961715, AA974649, W16617, AI092738, AW207722, AA233142, T64223, N79582, M27717, M73720, S40234, J05118, U67914, M73718, M73719
1857	HCROB02	877283		AI168748, AI376972, AI681157, AI279540, AA420977, N40163, AW AI274962, AW080693, N51345, AW337551, AA 440981, AA129415, AA AW367007, AL041883, AA720670, AA281119, D62242, R55623, AA83 1640690, AI695207, AA N23186, AW297680, AI N23185, AA843537, AI AW070934, D63021, AI
1858	HFKIN68	877284	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by	AI633741, AI017113, AA305124, AA227077, X58531

			the general formula of ath where a	
			inv integer between 1 to 1716 c	
			15 to 1730, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1858, and where b is greater	
			than or equal to a + 14.	
1859	HWHGC93	877285	Preferably excluded from the	AW275818, AI969511, W68529, AA627916, AW275825,
			present invention are one or more	W68815, AI375939, H42716, AI611676, R48249,
			polynucleotides comprising a	AA642987, AA631033, R73789, AI800001, AW452308,
			nucleotide sequence described by	3, AA730105
			the general formula of a-b, where a	AA933672, H25944, AI745535, AW276480, D29313,
			is any integer between 1 to 876 of	AW381131, AW380949, C00410, AW381579, AW381130,
			SEQ ID NO:1859, b is an integer of	AI220849, H25979, AA368136, AL035408
			15 to 890, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1859, and where b is greater	
			than or equal to a + 14.	
1860	H2CBC75	877287	Preferably excluded from the	
			present invention are one or more	D51060, C14389, C14014, D80522, F13647, D81026,
			polynucleotides comprising a	AW177440, Z21582, D81111, AW177501, AW177511,
			nucleotide sequence described by	C14227, D58283, D80043, T03116, D59610,
_			the general formula of a-b, where a	AA305578, D80022, C14331, D50979, AW369651,
			is any integer between 1 to 544 of	AW178986, D80168, D80247, AA285331, D51022,
			SEQ ID NO:1860, b is an integer of	AA514188, AA305409, D598
			15 to 558, where both a and b	D80195, D59467, D51423, D59619,
			correspond to the positions of	D80391, D80164, D59275,
			nucleotide residues shown in SEQ ID	D59787, D80227, D59502, D80439, D80241, D80014,
	•		NO:1860, and where b is greater	T11417, D81030, AW352117, D80188, D80269,
			than or equal to a + 14.	D80024, D80212, D80366, D80196, D59653, D80219,
				D57483, D59927, D80248, AA514186, D51103,
				C15076, D80064, D59889, D80193, C14429, T03269,
				AI557751, AW352120, D80045, D80133, D80378,
				D51759, D80302, AW178762, C14407, D80157,

AW377671, AW178893, AW177734, D80 D52291, AW178759, D59373, C75259,	AW378533, AW375405, AW360844, C14077, D59627, T02974, C06015, C14298, AW178906, H67866,), D51213, AW179328, C05695, AW	AW378539, AW360817, AW179020, T48593, AW378532, am378532,	1, AW179332, AW377672,	AW177731,	AW179024, AW377676, D45260, AW177505, AW178775,	λ,	AW178908, H67854, AW179018, AW178971, AW360834,	C05763, C14344, AW367950, AW179009, D60010,	AW179012, AW178980, AW178914, AW178774,	AW178781, AW177733, AW378543, D80258, H67858,	D59474, D58246, C14973, C14957, AI525917,	AI525227, D59317, D58101, D59503, D51221,	AW378525,	AA51418	i, D60214, AI525235, D59551,	AI525925, AI525215, Z33452,	D45273, AI525242, Z30160, AW378542, C13958,	237, AI905856, AI525222, T02868, A	D52317, D31458, AB002804, D86959,	J132110, AF058696, A62300, AB028	A84916, A62298, AR018138, A8259	5, AB002449, I50126, I50132, I5012	X68127,	Y17188, D26022, A25909, A45456, A26615,	AR052274, A94995, AR054175, Y12724, AR066488,	, AR03866	Y17187,	D50010, A632	AR016808, AR008408, AR025207, AR016691,
																								-						

				AR016690, U46128, A64136, A68321, AR060133, I79511, D13509, AF123263
1981	H2LAW79	877288	y excluded from the	,23, D80268, AA305578, D59502,
			present invention are one or more	D80164, D50979, C06015, C14389, D80038, F13647,
			polynucieotides compilsing a	332/3, D80133, AMI/8/33, D80188, 80227. AWI/8986, AA514188, D58283
			the general formula of a-b, where a	
			ny integer between 1 to 829	C14331, D80166, D50995, D51423, D59619, D80210,
			SEQ ID NO:1861, b is an integer of	D80240, D80253, D59787,
		,	15 to 843, where both a and b	D81030, D80378, D80212, D80193, D80196, D80219,
			correspond to the positions of	AA514186, D81111, AW378533, D59927, T03116,
			nucleotide residues shown in SEQ ID	D80045, D81026, D59610, D57483, C14227, D80439,
			NO:1861, and where b is greater	D80522, D59889, T03269, D80024, D80247,
_			a)	AW177440, D51103, D80248, D80241, D80366,
				D80302, C14014, Z21582, D59695, AW178893,
				D80133, AW178906, D52291, D80064, D80157,
				AW360811, D80168, D80014, AW375405, AW179332,
-				C14298, AW179328, D59503, AW178754, AW179019,
			-	AW378532, AI525923, AA809122, AW366296,
				D59317, AW
-				AW375406, AW377676, AW378534, AW352171, T48593,
				AW178762, AW179024, AW178971, C03092, AW378528,
				H67854, H67866, T11417, D59627, AW177456,
-				AW178907,
				AW360834, T02974
				D51221, C14973, AI535686, AW367950, AW178914,
				AW178774, AI525227, AW378543, D59474, AI525920,
				9, D31458, H67858, AI525925,
				C16955, C14077, AI525912, Z33452, AI525903,
				42, AI525215, Cl3958, AA305720
				T03048, Z86064, AL049679, AJ132110, A84916,

			1	
1862	HCE2C40	877289	preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 250 of SEQ ID NO:1862, b is an integer of 15 to 264, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1862, and where b is greater than or equal to a + 14.	, AF059650
1863	НМСDН54	877290	preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 1868 of SEQ ID NO:1863, b is an integer of 15 to 1882, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1863, and where b is greater than or equal to a + 14.	AL133778, AW408536, AA397575, AA399688, AA725429, AA324765, AA321795, AW243558, R86033, AW271180, H65207, AL134927, AB032995, AB018253

AW268628, AW408344, AI042425, AA286908, AI093993, AW316896, AI339306, AA736991, AI271364, AI539564, AA287969, AI689236, AI271364, AI539564, AA287969, AI689236, AI240770, AA035024, AA035512, AA804433, AW001846, AI191237, AI161031, AI015252, AW192454, AI817128, AI867530, AA557231, AI452866, AA804383, AL043242, AA627583, AA809613, T27814, M30818, M33883, AC004497	AW263526, AA457032, AW136358, AA828242, AA313271, AL078644	
Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 1912 of SEQ ID NO:1864, b is an integer of 15 to 1926, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1864, and where b is greater than or equal to a + 14.	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 544 of SEQ ID NO:1865, b is an integer of 15 to 558, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1865, and where b is greater than or equal to a + 14.	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 335 of SEQ ID NO:1866, b is an integer of 15 to 349, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1866, and where b is greater than or equal to a + 14.
877295	877298	817299
HTPFG64	H2CBQ45	нсQAD77
1864	1865	1866

1867	HKLSB60	877301	Preferably excluded from the	AA226684,
			present invention are one or more	AA002207, AA225124, AA225347
			polynucleotides comprising a	
			nucleotide sequence described by	
			wher	
			SEQ ID NO:1867, b is an integer of	
			15 to 536, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1867, and where b is greater	
			than or equal to a + 14.	
1868	HLHTC92	877310	Preferably excluded from the	R66025, R76969, AW043721, AA553904, AI417134,
			present invention are one or more	R58054, U77970, AR059959, U51625, U77969,
			polynucleotides comprising a	AR059960
			nucleotide sequence described by	
			the general formula of a-b, where a	
			is any integer between 1 to 839 of	
			SEO ID NO:1868, b is an integer of	
			15 to 853, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1868, and where b is greater	
			than or equal to a + 14.	
1869	HWLXP93	877319	Preferably excluded from the	AI968101, AI806911,
			present invention are one or more	AI918763, AW021370,
			polynucleotides comprising a	AA884471, W49632, T7
			nucleotide sequence described by	AW020878, AA812095, AA805395, AI767210, H08971,
			the general formula of a-b, where a	AA909382, AA325979, AA805574, AI911384,
		_	is any integer between 1 to 1232 of	AI520787, AC007239, U79290
			SEQ ID NO:1869, b is an integer of	
			15 to 1246, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1869, and where b is greater	
			than or equal to a + 14.	

AA299388	AC005037	
	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 408 of SEQ ID NO:1871, b is an integer of 15 to 422, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1871, and where b is greater than or equal to a + 14.	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 615 of SEQ ID NO:1872, b is an integer of 15 to 629, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1872, and where b is greater than or equal to a + 14.
877320	877321	877324
HUKBCSS	не9ғн60	ннеғс89
1870	1871	1872

1873	HCEOFOR	877326	Preferably excluded from the	N20930, AL135016, AL134824, AA702162, C03031,
))		present invention are one or more	', AI139490, AW057590
			polynucleotides comprising a	AI521171, N27797, AI953095, AI307324, AA705112,
			nucleotide sequence described by	AA969165, AA284734, AA325231, AI219990,
			the general formula of a-b, where a	AA287154, C03026, AI122656, AA772255, AA782094,
			iny integer betwe	AW073074, AI685711, AW192900, AI659385,
			SEQ ID NO:1873, b is an integer of	AI001129,
			7, where	AW296185, AA044143, AF034374, AJ224328
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1873, and where b is greater	
			than or equal to a + 14.	
1874	HLHBZ17	877327	Preferably excluded from the	C15947, H86703, AA359866, D61503
			present invention are one or more	
		•	polynucleotides comprising a	
			nucleotide sequence described by	
			the general formula of a-b, where a	
			is any integer between 1 to 693 of	
			SEQ ID NO:1874, b is an integer of	`
	-		15 to 707, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1874, and where b is greater	
			equal to a + 14.	
1875	HWLRP86	877329	Preferably excluded from the	_
			present invention are one or more	, AA394118,
			polynucleotides comprising a	AI870692, AI635237, AI139325, AI286284,
			nucleotide sequence described by	AW298025, AI830613, AA736608, AW008771,
			the general formula of a-b, where a	AW004643, AI277887, AI040732, AA628965, W93926,
			is any integer between 1 to 251 of	AI352001, AA954225, AI278572, N33931, AI128499,
			SEQ ID NO:1875, b is an integer of	W46369, AI159880, AI362660, AI350268, AA622742,
			15 to 265, where both a and b	AA887292, AI276858, AA250840, AA437277,
			correspond to the positions of	AA039774, AI242916, AI187707, AA804951,
			nucleotide residues shown in SEQ ID	AI277891, N63418, AA557131, AA662472, AI251864,
			NO:1875, and where b is greater	AI097294, AA991440, H99028, AI572652, AI610660,
			ω	AA055193, AI378407, AA719806, AI423797,

				AA729670. AA446337. AI311820. W81234. AI	AI300798,
				AA447436, AI189310,	
				2, AIS89143, AA918355,	
				AI095636, AA563972, N39264, N62211, AA936816,	6816,
				AA932784, AI868453, AW088157, AA970862, R7795	R77959,
				AI205800, N32013, AI582264, AI376345, AI22448	224485,
				AI274254, AI334251, AI401393, AI079459,	
				AI091021, AI277813, C14412, AI626008, AI279571,	279571,
				R26078, D80204, AA621068, AI400442, R80543,	143,
				AI479083, AA641535, AI378637, W81271, W8121	1215,
				R62807, H00547, C14369, AI784466, AI160567,	,67,
				AI160569, C14400, AI926459, C14352, AA44235	12355,
				C14220, C14335, AA687810, C14509, AA907451,	151,
				AW025906, AA459765, AL040127, AF125099,	
				AR029580, AF194030, AL133075, S7771, AF114784	114784,
				9, AL117443, AF207750, AL133645,	U67958,
				S78453, AL137554, Z30970	
1876	HISEQ81	877331	Preferably excluded from the	AA251070, AA663366, AL035663, AC008085,	U85196,
			present invention are one or more		
			polynucleotides comprising a	AL133247, AC004897, AL031390, AF135487,	Z83850,
			nucleotide sequence described by	, AL109922, AL034410,	
			the general formula of a-b, where a	, AP000500,	
				AC000064, AC007566, AL031775, AL023581,	
			SEQ ID NO:1876, b is an integer of	AC023172,	AL008629,
			15 to 513, where both a and b	AC004020,	AF072499,
			correspond to the positions of		24464,
			nucleotide residues shown in SEQ ID	AB024457, AB024458	
			NO:1876, and where b is greater	AB024479, AB024484, AB024488, AB024459,	
			than or equal to a + 14.	AB024469, AB024471, AB024478, AB024481,	,
				AB024462, AB024467, AB024463, AB024470,	
				AB024473, AB024475, AB024474, AB024482,	
				AB024476, AB024465	
1877	HWLWA0	877332	Preferably excluded from the	AA779795, AI808514, AA632293, AW263707,	
	7		present invention are one or more	, AI573067,	
			polynucleotides comprising a	, R38583, N6	97783,
			nucleotide sequence described by	AA889997, AW020741, AW084236, AI961833,	

			the general formula of a-b, where a	AW409834, AI914107, R37238, AI	AI202244, AW050863,
			teger between 1 to 636	AA318265, Z39970,	AI767672, AA757332,
			SEQ ID NO:1877, b is an integer of	AIS57697, AIS47137, T69960, AI	AI541216, AI535787,
			15 to 650, where both a and b	AI547038, AI557382, AI541533, AL122101	AL122101,
			correspond to the positions of	AL008582, AL035659, U44059, U0	U06935, Y11149,
			nucleotide residues shown in SEQ ID	AJ132931	
			NO:1877, and where b is greater		
			than or equal to a + 14.		
1878	H2CBS31	877333	Preferably excluded from the	AI248204, AA677184, AI380963,	AA284845,
			present invention are one or more	AW081587, T18597, AI525556, AI557084,	I557084, C14322,
			polynucleotides comprising a	AIS41205, AIS25500, AIS57533,	H65400, AW023216,
			nucleotide sequence described by	AIS57082, AA308485, AIS41321,	AI557731,
			the general formula of a-b, where a	AI557238, AI557263, AI557602,	T69960, AI541034,
			is any integer between 1 to 707 of		AI52585
			SEQ ID NO:1878, b is an integer of	AIS57543, AIS41027, AIS35994,	266121,
			15 to 721, where both a and b	A62298	
			d to the positi		
			nucleotide residues shown in SEQ ID		
			NO:1878, and where b is greater		-
			than or equal to a + 14.		
1879	H2CBN88	877334	Preferably excluded from the	l O	AI581828, A59459,
			present invention are one or more	A59517, AF048695, U52377, A59470, U53138	470, U53138,
			polynucleotides comprising a	A59468, U52375, A59469, U52376, A5946	6, A59466
			nucleotide sequence described by		
			the general formula of a-b, where a		
			is any integer between 1 to 550 of		
			SEQ ID NO:1879, b is an integer of		
			15 to 564, where both a and b		
			correspond to the positions of		
			യ		
			NO:1879, and where b is greater		
			than or equal to a + 14.		
1880	HWLOK01	877336	Preferably excluded from the	AI287235, AA587620, AA729307,	AI821703,
			present invention are one or more	AI688112, AI767799,	AA887822,
			polynucleotides comprising a	AA973956, AI693558, N78520, AI824444,	I824444, AI609594,
			nucleotide sequence described by	AI682837, AI690813, AI584118, AI824357	AI824357,

			the general formula of a-b, where a	AI224373,	AI886355,	AI537516,	AW167777,
			is any integer between 1 to 263 of	AI911020,	AI567802,	AW151451,	AI954293,
			:1880, b is an inte	AW194014,	AI888095,	AI439903,	AW079859,
				AI885905,	AI635528,	AI049669,	AI689096,
				AI636309,	AW131165,	AW090681,	AW084440,
			nucleotide residues shown in SEQ ID	AI538008,	AI784230,	AI491710,	AI925164,
			NO:1880, and where b is greater	AI220828,	AI432532,	AI696714,	AI472566,
			equal to a + 14.	AI874238,	AA761557,	AI251221,	AI620643,
				AI886940,	AI285439,	F34241, A	F34241, AI553926, AI628325,
				AI559863,	AI954095,	AA743430, AI804505	AI804505,
				AI357902,	R39624, A.	I918554, A	R39624, AI918554, AW079572, AW084896,
				AI580694,	U82987, AC	AC005218, I	I09499, AF109683,
				AL096728,	AJ001388,	X52220, U	AJ001388, X52220, U57715, AF188712,
				X95310, U	51123, AFO	81571, X66	X95310, U51123, AF081571, X66975, X57084,
				U79523, X6	56862, AFO	90923, ABO	X66862, AF090923, AB031064, X68560,
				AF078844,	AF114818,	122272, A	AF078844, AF114818, I22272, AL137663, E02253,
				X60786, A	7002672, M	92439, X99;	X60786, AF002672, M92439, X99226, X98066,
				AL133067,	AJ132433,	AJ132433, AF153205, AF167995,	AF167995,
				AR064250,	AF119337,	AL133069,	AL133069, AF114170,
				AF200464,	AF090886,	X63574, Y	Y08769, AR012379,
				AF141976,	X06146, Al	F077051, A	X06146, AF077051, AF003737, L40386,
				A65341, AL080146,		05032, ALO	J05032, AL050108, AJ012755,
				AF038847			
1881	H2CBR23	877338	Preferably excluded from the	AW340662,	AW316660,	AI970681,	AA889159,
			present invention are one or more	AI458059,	AI590367,	AI679607,	AI797703,
			polynucleotides comprising a	AW338264,	AI739401,	AA523715,	AA425084,
			nucleotide sequence described by	AI216290,	AA515788,	AA526334,	AI677745,
			the general formula of a-b, where a	AA134355,	AI674509,	AA143532,	AA313282,
			is any integer between 1 to 2508 of	AA927236,	AA315699,	AI620159,	AA922890,
			SEQ ID NO:1881, b is an integer of	AW062635,	AW374778,	AA100752,	AW374734,
			15 to 2522, where both a and b	AW368107,	AI214469,	AA134354,	AW368106,
			correspond to the positions of	AA385843,	AI919003,	AW379835,	AW389815,
			nucleotide residues shown in SEQ ID	AW206252,	AA213695,	AA305544,	AW418789,
			NO:1881, and where b is greater	AW368007,	AW368008,	AW374786,	AA313396,
			than or equal to a + 14.	AI940533,	AI940454,	AW062630,	AI920939, R25623,
				AW176592,	AA376950,	AW389787,	T48510, AW178927,

				, AW262708, AA626931,
				AI219498, AW390912,
				AW391129, AW379257, AW391053, AA746736,
				981, AW276
				٠.
				AF132818, D14520, AF079852, D82785
1882	HCYBK82	877339	Preferably excluded from the	, AI970681,
			inventic	AA425084, AW316660, AI458059, AI739401,
			polynucleotides comprising a	_
			nucleotide sequence described by	AI677745, AI216290, AA515788, AI674509,
			the general formula of a-b, where a	AA134355, AW338264, AI620159, AA100752,
				AA927236, AW206252, AI273521, AI919003,
			SEQ ID NO:1882, b is an integer of	AA626931, D59859, D80227, D80269, D80195,
			15 to 455, where both a and b	
			correspond to the positions of	u,
			nucleotide residues shown in SEQ ID	D59467, D51423,
			NO:1882, and where b is greater	9, D80378, D80210, D80240
			than or equal to a + 14.	
				D80219, D59927, D57483,
				C14389, D80241, C15076,
				9, D80045
			-	D51060, AW178893,
				51022, AW179328,
				AW378532, AW418789, AW369651, D80522, D58253,
				C14227, D80168, AW352158, D80251, D81111,
				6, D80248,
				AI910186,
				AW352117, AW360811, C14407,
				AW176467, AW375405,
				D80132, AA285331, AW366296, AW360817, AW375406,
				AW378534,
		_		
		_		T03116, AW360834, AW352172, AW352174, AW177505,
				AW178909,
				AW352170, AW177731, AW178754, AW179019,

			AW179018, AW179024, D59373, D80247, AW179220, AW179020, AI557751, AW177456, AW179329, AW178980, AW17733, AW378528, AW178908, AW178980, AW17733, AW378528, AW178908, AW179009, AW177733, AW378528, AW178017, AW179004, AW179009, AW179012, AW178914, D80114, AW367967, D801157, AW177022, AW178911, D51759, AW177728, D80157, AW177722, AW178911, D51759, AW17774, D80157, AW17722, AW178911, D51759, AW17774, D80258, AW177722, AW17723, D59627, T48593, AW378539, C14975, AW17723, D59627, T48593, AM378539, AI557774, AI535850, C14973, T02974, D45260, AW378533, H67866, AW367950, D51213, AW177508, AA809122, H67854, C03092, D80228, AW17734, AW17898, D59317, AI525227, D60214, T03048, AI525917, AI535686, C14344, D14520, AW17381, AM178986, D34614, D88547, A6298, AR018138, X67155, X17188, D26022, A25909, A67220, D89785, A78862, D34614, D88547, X19525, U87250, A86792, X93549, AR06488, AR016514, AR060138, A45456, A26615, AR06488, AR016514, AR060138, A4556, A26615, AR038669, AR066480, X1287, I18367, I18442, A30438, AF135125, D88507, AR038669, AR036697, AR038669, X17187,
			R008277, AR008281, AR008408 R016691, AR016690, U46128, 79511, A64136, A68321, AR06 AB023656, U87247, U79457,
1883 HCRMK82	K82 877340	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by	AK032065, X93535, AK008382 AW262592, AW367357, AI953876, AW265047, AI290247, AI261967, AA826909, AI336616, R46813, AA055350, R39815, N73560, H16260, AW365173, AC006251, X68487, M97759, AR044912, I20962

			1 2 2 2 1 1 2 2 2 1 2 2 2 2 2 2 2 2 2 2	
		_	is any integer between 1 to 844 of	-
			SEQ ID NO:1883, b is an integer of	
			15 to 858, where both a and b	
			correspond to the positions of	
	-		nucleotide residues shown in SEQ ID	
			NO:1883, and where b is greater	
			than or equal to a + 14.	
1884	HDTB006	877344	Preferably excluded from the	AI627846, AI686196, AI766030, AA159730,
			present invention are one or more	AA159731, AI478216, AI745281, AA683246,
			polynucleotides comprising a	AA252582, AW085579, AA936240, AA464699,
-			nucleotide sequence described by	AA732427, F11142, N62186, AA825887, N90846,
			the general formula of a-b, where a	N77132, AA376347, F08813, H50638, AL121257,
	•		is any integer between 1 to 1405 of	AL021937
			SEQ ID NO:1884, b is an integer of	
_			15 to 1419, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1884, and where b is greater	
			than or equal to a + 14.	
1885	HEGAM94	877346	Preferably excluded from the	AI935271, AI762915, AI809275, AA398950,
			present invention are one or more	AI127111, AI813351, AA749298, AA705921,
			polynucleotides comprising a	AA766587,
			nucleotide sequence described by	AI052069, AA291984, AA715043, AA460658,
			the general formula of a-b, where a	AA804876, N44967, AA394137, AW071467, N93279,
			is any integer between 1 to 1999 of	AI343843, AA393817, AI452856, AA292934, R90963,
	•		SEQ ID NO:1885, b is an integer of	W72279, AA861873, AA526081, AI819873, AA226137,
		_	15 to 2013, where both a and b	AA262543, R72676, T17354, AA514931, R73310,
_			correspond to the positions of	R90959, W25119, R64455, AI783605, W76306,
			nucleotide residues shown in SEQ ID	AI624523, AA490863, AA261906, AI864544,
			NO:1885, and where b is greater	
			than or equal to a + 14.	
		_		AI057127, R48640, R18641, AA461005, AA261923,
		_		R18640, H83702, Z38970, N36710, AL134185,
		_		H90736, H59529, H90786, AI784395, AA652150,
				AA652026, H60402, Z42828, AA226136, AA776284,

	TOTAL TOTAL TOTAL TOTAL	Alby6435,
	398313, R4	1473208,
	AI862134, AI273856, AL036705, AI539260	0,
	AI673140, AA715307, AA809974, AI369807	,7,
_	AL135047, AI440260, AW083572, AI554344	4,
	AA580663, AI683972, AI440238, AW151974	4,
	AI923989, AI440263, AI683568, AL138376	6,
	AI554821, AW020561, AA641818, AA761557	7,
	AA74835	3,
	AW055075, AI432644, AI538298, AI089748	8,
-	AI587000, AI590043, AL134830, AI682640	0,
	AI954080, AI691131, AI572396, AW087262	2,
	AI094749, AW162194, AI613038, AI557104	4,
	, AI47527	٥,
	AW087445, AI625293, AA065052, AI289310	0,
	AI678857, AI445505, AI370965, AA282824	4,
	AI866457, AI872423, AL135012, AI591093	3,
	AI56758	2,
	AI53880	5,
	AW07538	2,
-	, AL119748, AI91579	5,
	AW243886, AW130129, AI925736, AW168012	2,
	AI798114, AL121270, AA609644, AI440236,	6,
	AW268122, AI680221, AI064830, AI473471	1,
	9, AI283322, Y11254, AR050959,	AC002464,
	AJ238617, AF150103,	D44497,
		93914,
		1,
		37539,
	AL137459, AF082526, E12888, AF145233,	AF118094,
	AL133113, U92992, AC002287, AF017437,	133391,
	AL133637, AF069506, AL122101, AL133080	0,
	AL133053, AL122049, U70981, AF115392,	
	, U67082, AL137284, AR03482	X1513
-	, AL137479, AF051325,	\$63521,
	AF004162, AF161413, AJ238093, AL122110	0,

-				ALO80074, ARO66486, E12580, ALO50149, U51123, AP146568 H53505 AP064250 V10655 AL137526
1886	HDTAH72	877347	Preferably excluded from the	1
			present invention are one or more	
			polynucleotides comprising a	AA621945, H97851, AW082375, R34105, AA376468,
			nucleotide sequence described by	AA376668, AA376330, AA224458, R34106, AA166983,
			w	D58161, AI919577, C21057
			is any integer between 1 to 1879 of	
			SEQ ID NO:1886, b is an integer of	
			15 to 1893, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1886, and where b is greater	
			equal to a + 14.	
1887	HARAG42	877351	Preferably excluded from the	AA534438, AA296922, AI732343, AA502919,
			present invention are one or more	AI732203, E13091, AR028526, AF048700, E13090
			nucleotide sequence described by	
			the general formula of a-b, where a	
			is any integer between 1 to 419 of	
			SEQ ID NO:1887, b is an integer of	
			15 to 433, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1887, and where b is greater	
			than or equal to a + 14.	
1888	HCQDL20	877355	Preferably excluded from the	R10554, AA873089, AW007836, AA376913, AA702706,
			present invention are one or more	AI861809, AI052145, N74374, AI739300, AW055276,
			polynucleotides comprising a	T40120, AA343939, T40984, J04813, AF209389,
			nucleotide sequence described by	S53047, M14096, M18907, X12387, J04449,
			the general formula of a-b, where a	AF182273, D31921, Mi3785, X90579, L26985
			is any integer between 1 to 399 of	

			SEQ ID NO:1888, b is an integer of	
			15 to 413, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1888, and where b is greater	
			than or equal to a + 14.	
1889	HLQGF34	877356	Preferably excluded from the	
	,		present invention are one or more	
			polynucleotides comprising a	T98255, N74426, AA376913, AA416822, T40120,
			nucleotide sequence described by	AI861809, AI678780, AA343939, T98311, AA878869,
			the general formula of a-b, where a	AI761228, X90579, L26985, AF209389, J04813,
			is any integer between 1 to 769 of	S53047, X12387, M14096, M18907, J04449, D31921,
			SEQ ID NO:1889, b is an integer of	AF182273, M13785
			15 to 783, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1889, and where b is greater	
1890	HCDCF78	877358	Preferably excluded from the	AI703276, AW188039, AA451771, AA316434,
			present invention are one or more	AI690259, AI681353, AA045904, T29610, AI627945,
			polynucleotides comprising a	AW188125, AW188144, AA099043, AW237788,
			nucleotide sequence described by	AI470110, AW170058, AI654577, N21480, AI678192,
			the general formula of a-b, where a	AI745496, AW292165, AA449964, AI167571,
			is any integer between 1 to 385 of	AI186510, AI392894, AI459190, AW196865,
			SEQ-ID NO:1890, b is an integer of	AI761196, AI199686, AA767664, AW373992,
			15 to 399, where both a and b	AI129612, AI272655, AI272824, AW051688,
			correspond to the positions of	AI765956, AI220043, AA099044, AI681033,
			nucleotide residues shown in SEQ ID	AI628056, D17400, M97655, D25234, L76259,
			NO:1890, and where b is greater	M77850, U63380, U63381, U63382, U63383
1891	HMIBE59	877361		AL043108, AI912625, AI268389, AA541465,
			present invention are one or more	AA626702, AI814451, AA703936, AW137200,
			polynucleotides comprising a	AI769406, AI814300, AA843784, AI677825, N90942,
			nucleotide sequence described by	AL133947, AI122639, AI583230, AI956122, W58349,
			the general formula of a-b, where a	AA043151, AI911861, AI146802, AA433844,
			is any integer between 1 to 3021 of	AA829527, AI829684, AA393149, AI248810,

AI925678, AA062558	0 - 0	AW403227, AA644436, N35139	8100, 6227		1, R23998, AIS97694,	R24938,	T28111,	A1525244, AA042803,				AA098979,	811590,		00000	AA329386, 3906,	H00588,	100143,	D31543,	3940,	AA069494,	AI370594,	67,	,
9, A16343 2, W48807 W52177, AA639344	AI08857 6, W5847	N36852, AW440100, AA708923, AW AA846487, AI075216, N56895, AA	AWO80740, N46123, AP R80330	AI185045, AW204631,	, AA305934, AA158097, AW027841 AA262561, AA626808, AA040760,	. ∝	AA165152,	AA169476, 2, N32999,		AA65412	A1435866, AA478972 A1683540, A1242454	098811, AI970953,	N24550, AI656583, AA098926, AI811590	AA771762,	AW300195	, DZUIU4, AMBIU/U5, AM323386 AI824554, T70014, R23906,		2731, AI		59266, R23125, H839		T69942, AA3197		AI597664, AI972622
AW148927, A1693209, AA31332 AA165311, AW015279, AA43556 N33995, AW337556, AI200909, C75536, AA740996, AT056139.		AI090980, N36852, AW AA746255, AA846487, W20313 W20170 W202	AAO63056, MJZ178, MUOZ AAO63056, C75383, AW AARRASC, AT339843	R23907, AW272245, AI	AI347721, AA305934, AA158097, AW027841, N36871, AA262561, AA626808, AA040760, A3		N46141, AA165180, H94816,	, 120138, Ar , AA366030,	_		A14/0822, A1559820, A1672499. AA782245.		~		AIS65050, AI669676,	, AA991913 , H80964,	F00987,	R45201,	AI681692, AI015103,	3		AI674511	R23126,	AI420216, AI365551,
SEQ ID NO:1891, b is an integer of 15 to 3035, where both a and b correspond to the positions of nucleotide residues shown in SEO ID	and where b is greater qual to a + 14.																							
													-											

				AA243213, T35681, C04078, C75653, T11331,
				W23989, T18555, T11401, T39150, AA094342,
				AI824772, W17101, N91885, AA453560, T11352,
				T10404, N47782, AA091310, C00888, AA165310,
				T27528, AA248615, AI420657, R79019, T25720,
				AA863104, AI095737, T11400, AA523550, AA913502,
				AI218901, AI827982, A93912, M31470, A93910,
				ഗ
				AL035361, R62747, AA853568, AA916254, AA969277
1892	HMKAK86	877363	Preferably excluded from the	AA190594, T40630, AI920974, AI055924, AW081296,
			present invention are one or more	
			polynucleotides comprising a	AA053866, AI923333, AA516448, AA344620,
			nucleotide sequence described by	AA347824, H05424, H02246, R22341, T40694,
			the general formula of a-b, where a	AA344748, AW449318, AA737586, AI950008,
			teger between	AA037725, AA345669, AA302793, AA302797,
			SEQ ID NO:1892, b is an integer of	AI355125, T39494, AW150691, AA902521, AI278972,
			15 to 376, where both a and b	AI270407, AB033054
			d to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1892, and where b is greater	
			than or equal to a + 14.	
1893	H6EDF71	877370	Preferably excluded from the	AI961479, AI923277,
			present invention are one or more	AW129387, AI432621, AI701980, AI613026,
			polynucleotides comprising a	н
			nucleotide sequence described by	AI280720, AA918056, AA938271, AA418701,
			the general formula of a-b, where a	AI338213, AI707674, AI476785, AA478755,
			is any integer between 1 to 1290 of	AI082024, AA455447, AA834685, AI742309,
		-	SEQ ID NO:1893, b is an integer of	AI857345, AW090377, AI708271, AI016116,
			15 to 1304, where both a and b	AA588253, AI167998, AI445021, AA455448,
			correspond to the positions of	AA669129, AI474588, AI208596, AW015585,
			nucleotide residues shown in SEQ ID	AW015582, AI283110, AA773711, AA558268, W93910,
			NO:1893, and where b is greater	D54259, W52496, AW195549, AA418855, AA937302,
			equal to a + 14.	AA960793, AA976090, AW105521, N62182, AA009747,
				AI686709, AW178327, AI275229, T39172, AA471190,

AI884496, T40454, AI889115,	AA773707,	H16423, X69398,	Z25524, D87659	AA861203,	AI587088,	AI818020,	AW190795,	AW192746,	AW190516,	AW192636,	AI683156,	AI685181,	AI627454,	AI984752,	AW074064,	AI697355,	AI587134,	AI813449,	AI432646,	AI913951,	AI571989,	AI587043, W94653,	AI190373,	AI250818,	AI992004,	AA921724,	N24418, AA622296,	AI074992,	AW337830,	AA854050,	W45594, AI610384,	N40742, AI076955,	AA904719,	AI280126,
AI88 T40			22524							AW1					AW0			AI8			Ċ									Ċ				AI28
AA535376, AW276245,	AW393132,	AA648104,		AW190863,	AI955634,	AW079778,	AI955860,	AW190680,	AI823711,	AI624269,	AI538927,	AI683833,	AI587424,	AI623652,	AI110775,	AI804583,	AI884376,	AI560022,	AI753639,	AI628183,	AI190931,	AW152597,	AI492736,	AI285408,	AW337268,	AA872416,	AI289514,	AI751083,	AA173912,	AI632052,	AI086679,	AW241380,	AI358461,	AI247519,
AW166867, AA969906,	AA279095,	AI932456,	AF017437, AB012693,	AI379830,		AI572602,	AI963206,	AI924265,	AW337223,	AI674875,	AI620393,	AW074297,	AW173674,	AW131016,	AI802264,	AI097497,	AIS70335,	AA910529,	AI333407,	AI818473,	AI587385,	AI198766,	AI868031,	AI971361,		AI191817,	AI962031,	AI753534,	AI559198,	AI304733,	AI751084,	AA716327,	AI754958,	AA947025,
AA777967, AA953028,	AW137558,	AW393156,	Z25521, A	AI625476,	AI952079,	AI926590,	AI978757,	AI587161,	AW152121,	AI623641,	AI573153,	AI860782,	AI923388,	AI453249,	AI084796,	AI571619,	AI445032,	AI754165,	AI028123,	AI683000,	AI193030,	AI520669,	AI520755,	AI571651,	AI299640,	AI313475,	AI754230,	AW152146,	AI436436,	AA722578,	AI680348,	AI086711,	AI692374,	AI262790,
				Preferably excluded from the	present invention are one or more	polynucleotides comprising a	nucleotide sequence described by	the general formula of a-b, where a	is any integer between 1 to 2603 of	SEQ ID NO:1894, b is an integer of	15 to 2617, where both a and b	correspond to the positions of	nucleotide residues shown in SEQ ID	NO:1894, and where b is greater	than or equal to a + 14.																			
				877373																														
				HOELCIS	~- -	-																		··· ,		_	_	_					<u></u>	
				1894																														

DIB30039 H96641 W76543 DIB19930	0. N31417.
TOUR STACK	849 AT752417
 AI582458	28732, C75417,
646, AI032902,	71822,
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	602, AA669993,
	02375,
AW339078,	75137,
AI439371, AW239521, AA887673, N2	AA887673, N24118, AI362463,
H99628,	215, AI815228,
	AI050040,
	84028,
AA076641, AA470703, AI689178, AI436443,	36443,
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\vdash	5, AA703635,
 AI862948, AW449712, AI579942, W04328,	328, AA962252,
	967, AW193168,
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, R00074, AW021966,	4, AA157265,
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AI131364, R66674,	W84537, AA661834,
, AI868207, AA642245	75185,
0, AW243595, R92565,	AI476033, AW198023,
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F085482,	S56205, M33300,
X81581,	791, AR018793,
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AB002449, AR008443, IS0126, IS0132	2, IS0128,

				I50133, AB012117, Y17187, I09494, A45456,
				AR066488, AR016514, AR060138, A26615, AR052274,
				AR008277, AR008281, A85396, AR066482, A44171,
				X64588, Y09669, A85477, A43192, A43190,
				X93549, U46128, AR066490, I14842, D88507,
				AR016691, AR016690, AR054175, D50010, I18367,
				A63261, AL133015, AR008408, I79511, AR062872,
				A68321, AR060133, A08456, A31057, T47722,
				3,
				, T97068, T98840, T99143,
				R21264, R31911, R31957, R62970, R63024, R63509,
	-			
				H79278, H79389, H85490, H96640, N20906, N30033,
				N31502, N74163, AA026408, AA040602, AA040685,
				AA1735
	_			D78982,
				AA852355
				F05444, AI360546, AI473496
1895 HAJ	HAJBN08	877375	Preferably excluded from the	AA350728, AA316351, AA112015, AA216692,
_			present invention are one or more	AW246040, AA693635, AW407512, N55660, AI362985,
			polynucleotides comprising a	AJ002190, AF043937
	•		nucleotide sequence described by	
			the general formula of a-b, where a	
			is any integer between 1 to 536 of	
			SEQ ID NO:1895, b is an integer of	
			15 to 550, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1895, and where b is greater	
\dashv	٠		than or equal to a + 14.	
1896 HFV	HFVHT62	877377	Preferably excluded from the	AI739135, AI066521, AW173105, AW261971,
			present invention are one or more	AL039012, AI954494, AA830348, AA284072,

			polynucleotides comprising a	AA789097,	AI005313, A	AA777794, AI041134,
			nucleotide sequence described by	AA856987,		AA769862, AA804528,
			ä	AA831168,	AA494334, A	AI143496, AI141222,
			is any integer between 1 to 843 of	AI372907,	AA831166, N	
			SEQ ID NO:1896, b is an integer of	AI075136,	AI076701, A	AA305065, AI076409,
			15 to 857, where both a and b	AA315766,	AI273523, A	AA450169, AA314707,
			correspond to the positions of	AA284166,	AA158102, A	AA158102, AI352491, AA257019, T96666
			nucleotide residues shown in SEQ ID	T28941, A	4352693, AA6	T28941, AA352693, AA627383, AA257103, AA464156,
			NO:1896, and where b is greater	AI206700,	T96781, AA1	T96781, AA158059, AA055005, AA757304
				AW059834,	AW340182, A	AW340182, AA092745, AI678081,
				AW368066,	L27711, U02	L27711, U02681, I30245, L25876,
				AL049778		
1897	HILBZ32	877378	Preferably excluded from the	AI739135,	AW173105, A	AI066521, AW261971,
			present invention are one or more	AI954494,	AA830348, A	AA789097, AA284072,
			polynucleotides comprising a	AA804528,	AI005313, A	AA777794, AI041134,
			nucleotide sequence described by	AA856987,	AI700317, A	AA831168, AA769862,
			the general formula of a-b, where a	AL039012,	AA494334, A	AI143496, AI141222,
			is any integer between 1 to 765 of	AI372907,	AA831166, A	AA769007, N64843, AI075136
			SEQ ID NO:1897, b is an integer of	AI076701,	AI273523, A	AI076409, AA305065,
			15 to 779, where both a and b	AA450169,	N92087, AA3	N92087, AA315766, AA158102, AI352491
			correspond to the positions of	AA314707,	AA257019, T	T28941, T96666, AA627383,
			nucleotide residues shown in SEQ ID	AA464156,	AI206700, A	AA257103, AA284166, T96781
			NO:1897, and where b is greater	AA158059,	AA352693, A	AA055005, AA757304,
			equal to a + 14.	AW059834,	AW340182, A	AI678081, AW368066,
		-		AA450104,	AA092745, L	L27711, U02681, L25876,
				I30245, A	AL049778	
1898	HAPOR25	877380	Preferably excluded from the	AW272420,	AW242297, A	AW242297, AA165082, AW263065,
			present invention are one or more	AI378393,	N34290, AA4	N34290, AA488409, AI347346, AA701568,
			polynucleotides comprising a	AI174216,		AI918787, AA948264,
			nucleotide sequence described by	AA594684,	AW299275, A	AI222510, AI243187,
			the general formula of a-b, where a	AW070414,	AI076437, A	AA488545, AA470051,
			is any integer between 1 to 3296 of	AW380452,	AA164540, A	AI076271, AA657436, N75339,
			SEQ ID NO:1898, b is an integer of	AI473793,	AW025483, AA701579,	A701579, N58947, AA577451,
			15 to 3310, where both a and b	R77252, A	A897628, T62	R77252, AA897628, T62571, AA102397, R77251,
			correspond to the positions of	AA704389,		AI697267, AA826647, W90783, AA632480,
			nucleotide residues shown in SEQ ID	AI032244,	AA583140, W	W01846, T31054, Z43387,

			NO:1898, and where b is greater than or equal to a + 14.	1, AI244271 T62961, AW	56, AA916276, AI0, D25970, N48191,
				AA252955, AA939180,	
				R06301, AW304307, R682	R68203, AW368013, AW364400,
				ALZ04114, AW364338,	N44181,
				1565221,	
				D12170, AW294181, T248	AW337772,
				N53338, W90688, AA253123,	7
				AI344295, AW364396, XT	X73882, Y15197, AL023284
1899	HELBN30	877384	Preferably excluded from the	1	AA654731, AA278203,
			present invention are one or more		AA057712, AI628148,
	,		polynucleotides comprising a	AI479111,	AI248082, W49737, AA009479,
			nucleotide sequence described by	AW449837, AA447481, R(R06619, AA040474, AI925539,
			the general formula of a-b, where a	AI347058, AA740520, WE	W86694, T29489, AA341731,
			is any integer between 1 to 1170 of	N59177, AA632345, AA057395	
			SEQ ID NO:1899, b is an integer of	AI805718, AA120879, HS	HS9542, AI379485, R25939,
			15 to 1184, where both a and b	Æ	
			correspond to the positions of	AA262292, AI425046, R(R01630, T50780, AA993907,
			residues	2, AI911765	, AA740339, AI186344,
			NO:1899, and where b is greater	AI583330, W25428, AI1	AI193756, AA001910, N75914,
			than or equal to a + 14.	AA921773, AW363532, AA	AA693648, AI242044,
				AI753406, AA588342, M	M60618, AF056322, U36501
1900	HHFMH12	877387	Preferably excluded from the	7, AI750041,	AI589918, AI971206,
			present invention are one or more	5, AI870013,	ω,
			polynucleotides comprising a	3, AW068564,	, AI43191
			nucleotide sequence described by	, AI422826,	AI493768, AI363488,
			the general formula of a-b, where a	_	AA100840, AI755276,
			is any integer between 1 to 3864 of	AA476207, AI992015, AV	AW026405, AI190217,
			SEQ ID NO:1900, b is an integer of	AI738539, AI439206, A <i>I</i>	AA037160, AI361483,
			15 to 3878, where both a and b	AA877117, AA425180, A	AI372673, D80801, AA678831,
			correspond to the positions of	AI376927, AA160849, A	N77542,
			nucleotide residues shown in SEQ ID	, AI084962,	AI356122, W88956, AI499098,
			NO:1900, and where b is greater	AA325211, N62261, N947	N94717, AA043409, AA789304,
			than or equal to a + 14.	AA355373, AI372674, He	H63354, AA313505, AA351821,

Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where is any integer between 1 to 161 of SEQ ID NO:1901, b is an integer of 15 to 175, where both a and b correspond to the positions of nucleotide residues shown in SEQ NO:1901, and where b is greater than or equal to a + 14. Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where is any integer between 1 to 1793 of SEQ ID NO:1902, b is an integer of 15 to 1807, where both a and b correspond to the positions of nucleotide sequence both a and b	HBXAC19 877388 Preferably excluded from the polynucleotides comprising nucleotide sequence described and its any integer between 1 to 175, where both a such polynucleotide residues shown nucleotide residues shown nucleotide residues shown nucleotide residued from the polynucleotides comprising nucleotide sequence described is any integer between 1 to 1850 ID NO:1902, b is an integer between 1 to 1850 ID NO:1902 ID NO
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		than or equal to a + 14.	AI498762,	AA865546, AI189894, A	AA740394,
			AA133324,	AI129125, AW022772, A	AA493572,
			AI202523,	AA676968, AA329249, W	W05485, AI038788,
			AA716709,	Ą	330025, AA126339,
			AI358727,	N56854, AA978006, AI7	AI719099, W37534,
			AA953629,	AA663651, AI693987, AA076372	A076372,
			AW090432,	N32431, AI362222, AA6	AA617762, AA782855,
			AI161045,		AI371415, AA136072,
			AW044060,	AA416713,	AI500608,
			AA991563,	6, AA305695,	1358972,
			AI926596,	AA384023, T40849, AA076501,	76501, AI991793,
			AA730185,	AI698869, AI949134, AJ	AA687665,
			AA121023,		AW275473,
			AA339483,	N35481,	AI363884, AA369524,
•			AA355468,	AA845483, F29460, W52	535, AI810861,
-			AA582099,	H19093, N80825, AA708946, AA384975	946, AA384975,
			AA379550,	AA373476, AA648147, AI818027	1818027,
			AA534415,	N56694, AW083204, AA372060,	72060, AA496767,
			AW007697,	AA748067, AI655704, AJ	
			AA042892,	M62297, AA043512, AA043513,	43513, AA384593,
			AA372059,	, AI279119,	AI635811,
			AA384973,	6, AA480294,	AI276970,
			AA515682,	AA043019, AA773750, AJ	AA169816,
			AL038644,	, AW080380,	AI434682,
			AA384974,	, AA176343,	AI278392,
			AA706110,	AA678943, AA515683, N2	N20394, AA375542,
			AR030958,	2, AC004922,	S77329, Ull861,
+			AF058791,	T39861, AI421422	
1903 HWHQH17	877393	7	AI346901,	AI191444, AW001394, AI	AL036955,
		present invention are one or more	AI660571,	AI818120, AI018511, A1	AI052368,
		ides comp	AW027921,	AW007170, AA603096, AM	AW057755,
		ednence	AA485948,	AI149233, AW081475, AJ	AI677997,
		b, where	AW410351,	AW300638, AA488667, AV	AW409854,
		teger between 1 to 2796	AA402239,	, AA486050,	AW409878,
		NO:1903, b is an integ	AA486507,	6, AW194332,	AA554501,
		15 to 2810, where both a and b	AW084623,	AW409835, AA617980, AI	AI040998,

		•		
correspond to the positions of	AI804511,	AW410178,	A1434575,	AI589609,
e residue	AA664262,	AW409614,	AA430234,	AA479644,
NO:1903, and where b is greater	AA488187,	AW305031,	AA410912,	AI313158,
equal to a + 14.	AA488684,	AI355319,	AA430559,	AI190998,
	AA676466,	AW409596,	AA476902,	AA878887,
	AA902228,	AI687559,	AI074371,	T51288, AA459629
	AW303926,	AA599915,	AA485902,	AI126733,
	AI445068,	AW409577,	AA593873,	AI016575,
	AA719627,	AA488240,	AA482604,	AW303900,
	AA486198,	AA430025,	AA847289,	AA188216,
	AW409876,	AI246054,	AA402700,	AA421202,
	AA416583,	AA847234,	AA630648,	AI802458,
 -	AA211469,	AA190840,	AW025006,	AA035463,
	AA186363,	AA992133,	AA670258,	AI469676,
	AA426620,	AA179226,	AW300817,	AI161092,
	AI199582,	AI339697,	AA993589,	AI083639,
	AW001456,	AA758347,	AA633544,	AA987682,
	AA486304,	AI889937,	AI581339,	W45576, AA701272
	AI565866,	AI347560,	AI079926,	AI146534,
	AA601655,	AI459359,	AA489322,	AI247541,
	AI469729,	AI074396,	AW001571,	AA579941,
	AI278644,	AI459387,	AA513381,	AA477332,
 -	AI076715,	AA976943,	AA833630,	AA149959,
	AI921791,	AI280849,	AI174208,	AI066715,
	AI285157,	וח	AA132930,	m
	AI269574,	H16257, AA	A588880, AA133075,	A133075, AA188878
 	AA627878,	AA025145,	AI568930,	AA196286,
	AI220665,	AA723359,	AA954162,	AA489559,
	AA630299,	AA135404,	AA188819,	AI362548,
	0	m		AI453521, AA804703
	H05127, A	AA477015, A	AI802650, T	
	AA665815,	AA186894,	AI984554,	AI984554, AA488648, W72251,
	AI094464,		W03180,	AA026596, AA112256
	AA486030,	H39838, A	[074194,	Н
	AW247688,	AA029620,	AI091141,	AI700362, H39837
	AA724925,	W69320, R	R76662, H95	672, W37885,

	H95068, AA612954, W37947, AA580556, H20424,
	70, H49118, H22277, AA120
	AA635599, AI032213, AI377
	A180255, AA046132, AI613018,
	, AI
	226,
	47, AI
	AI355337, AA654907, F07217,
	AA190498, H26731, AW166037,
	W15177, TS
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_	L00084, U37439, K01846, AC004616, K01845,
	9, Z23142, S69407, X77952,
	7442, Z36810, D16853, K
	3, L00082, L00083, M31698,
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	, T59510, T59556,
	T69436, T69569, T69637, T70491, T71461, T71584,
-	, T97732, T97836, R18156, R37533,
	, R41703,
	, H26732, H41805,
	, R92705, H49054, H51452,
	, H58770, H58822,
	, H79709, H90004,
_	N69990, N74472,
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	AA026595,
	AA128188,
	44, AA468336, AA503585
	, F15917, AA631927,
	88002, AA864500,
	AA482538, AA628208, AA69415, AA719284,

			8, AA852209, T10360, T1 AA694056, AI269768, AI 6, AI659249
	877396	y excluded nvention ar otides comp e sequence al formula teger betwee 1904, b is 9, where be d to the pod to the pod and where begund to a + iqual to a +	3, A1638226, AW014789, 3, A1075890, AW242842, 0, AW372249, AA630413, 2, AA134046, N32561, AI 9, AA551242, AA480899, 8, AA210774, W00846, AI 3, A1830594, AI589236, 5, AA489659, AI027334, AA923540, AA669903, W7 8, AI041901, AA126268, AA232572, AW002525, W0 AA700807, AA134045, AA 3, AA064885, AI093714, 5, AA902590, W46161, N2 5, AA902590, W46161, N2 6, AA664808, AA819060, AI675041, AI590268, R8 AA480245, AA991447, R8 AA480245, AA373949, R142 AW181920, AA374575, H0 R14352, AA373949, R142 1, N89241, R91989, N682 9, N72476, AI547027, AA 9, AW382356, AW371061, D50917
HCFMY07	877406	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 3975 of	AW004054, AL135021, AW173336, AA846316, AI208817, AA861115, AW377287, AI884576, AA403122, AW377237, AA449008, N22548, AI612907, AI697252, AI337225, AA488782, AA166884, AA114179, AA824590, AA723930, AA488998, AA534667, AI335733, AA922029, AA846011,

AA732053, AA807156, N31650, D61907, AA604009, AL121217, C75317, AI183839, AA285257, AI631612, AI701860, AI872948, AA724511, AA593781, AA03450358, AA348286, AW014127, AA034503, AW382984, AA114216, AA714035, N44341, AA034503, AW382984, AA114216, AA714035, N44341, AA0342769, AI695226, AA039307, D82808, T57805, AI865947, AA490260, D79331, H45236, AA312976, AI904624, R62919, D59331, H67517, R62920, T96420, R21224, D62945, AI648439, AW383006, AA789111, R63601, D62711, AA336494, AA340489, T37263, AI625255, H68430, AI824522, D82698, AZ1223, AI61720, N59296, AA249438, AI217233, D82710, D59332, AA565565, AA450364, R95490, AA490906, C01268, AW363022, AA913585, AA491092, E13124, U42424, U58512, U61266	AI963125, AI609225, AI884581, AW069271, AI953978, AI567519, AA703985, AI858101, AI281477, AA878466, AW084603, AA004204, AI755045, AI753615, AA122291, AW150834, AL038513, AA706823, AI814914, AA127736, N32519, AA706805, AI564735, AI670785, AI754803, AI160667, AI755281, AI122842, AI127349, AW088731, AI083555, AA609330, AA058930, AA486379, AW021109, W93848, AA115524, AI090089, AI570898, AI262822, AA903134, AI697486, AI698658, AA121511, AI580763, AL038512, AM439391, AI341677, W52306, AA010309, AW069115, AI127946, AI692736, AA600038, AW069432, AI754320, AI346302, AA723122, AA010310, AA599273, AA137194, AA599504, AW069432, AN088383, AI751005, AA725207, AW385359,
SEQ ID NO:1905, b is an integer of A. 15 to 3989, where both a and b. A. correspond to the positions of A. nucleotide residues shown in SEQ ID A. NO:1905, and where b is greater A. than or equal to a + 14. The positions of A.	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a his any integer between 1 to 2615 of SEQ ID NO:1906, b is an integer of 15 to 2629, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID A nucleotide residues shown in SEQ ID A than or equal to a + 14.
	877408
	HSYBP46
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	AW372817, AI446310, AW074603, AI075140,
	N89578,
-	AW372828,
	AA099729, N42734, W02000, AW372820, AI160542,
	AI095555, AI754231, AA070970, AW302579,
	AA114947, AI357733, AW386363, AI864906,
	2, AW393341,
	, AI750527, AA305175, AI200515,
	AI342335, AI751983, AI417127, AA993150, R77205
	AI077562, N75508, W93869, AI127162, AI582477,
	AW068212, AW393339, AW393324, W87515, AI751004
	AI039775, R95826, AA040410, N68613, AI752199,
	AA150616, AI919268, AW372823, W87487, AW393333
	AW088208, N43019, R95777, W30698, AW393343,
	AI671130, AI094661, R69515, AA330038, AA705256,
	AW393342,
	AA974667, N99050, AW068455, AI147454, H87987,
	62061, AA578679, AA
	AW393330,
	AI589497,
	σ.
	09536, AF
	AW235794, T27809, H0
	AA115948, AA853107,
	AA194797, AI263967,
	AA342316, AI569315,
	AA092962, AI750253,
	AA332339, AW196741, AI537624, AA040329,
	W57799,
	C02028, AA
	AA334576
	49492, T3
	87, T31612, R07858, AA332886, AA
	AA449254 COOO44 AA348035 AA328980 AA361011

	M11718, Y14690, X04758,
-	M10956, J03051, Y1
	1, D83032, AF106862,
	U49434, E02349, AL137550, AF137367, AF078844,
	3, AL133014
	214, AR038854, S7
	78525, AF113019,
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	049382, AL137558
	AL137463,
	AL050116, AL137555, AL137480, AJ00
	5, A08907,
	F067790, I89931,
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	A08908, U53505, AL13
	S76508, AL080163, AL080124, Z13966, I89934,
	AL050170, E02152,
	AB007812, U35846,
	AL133665, X79812, I03321,
	F017437, U67958, AL122106,
	AL117585, AL023657, AF199027, AL137548,
	AL133568,
	1, AL13757
-	AL050092,
	L137292, I6
	526, AL133054
	AF200464, AL117578,
-	AF185576, AL117629, S69510, AF055917, AF159615
	F162270, A1
	7, AL049300,
	H13204, H88165, H88165, N6428
	61456, AA594297
	AI086998, T03859, T24745, AI128830, AI537635

1907	HCRQK59	877411	Preferably excluded from the present invention are one or more	AI394016, AA503225,	AI337333, AW008484, AI492226, AI832480, AA551754, AW263863,
			polynucieotides comprising a nucleotide sequence described by	AA/825/3, AI631409,	AA4630/1, A1/00423, A1380330, W95477, A1651800, AA804581, AW016198,
				AI567909,	W05729, AW338263, AA488420, AW134932,
			is any integer between 1 to 1537 of	w	AI424300, AI569012, AA348345, W95367,
			SEQ ID NO:1907, b is an integer of	N74885, Z2	Z20694, AI569356, AW083000, AA745423,
			15 to 1551, where both a and b	35,	T24482, AI355870, R65920, AW054656,
			correspond to the positions of	A75401	
			nucleotide residues shown in SEQ ID		
			NO:1907, and where b is greater		
			than or equal to a + 14.		
1908	HWLXK44	877437	Preferably excluded from the	H53943, RC	R09272, W52643, AW001226, AI827422,
			present invention are one or more	AI086839,	AI086839, AI752330, AI752329, H53944, AL136295,
			polynucleotides comprising a	U94831	
			nucleotide sequence described by		
			the general formula of a-b, where a		
			is any integer between 1 to 454 of		
			SEQ ID NO:1908, b is an integer of		
			15 to 468, where both a and b	-	
		•	correspond to the positions of		
			nucleotide residues shown in SEQ ID	,	
			NO:1908, and where b is greater		
			than or equal to a + 14.		
1909	HE8DZ94	877630	Preferably excluded from the	AI684587,	AA610052, AI189791, AI186697,
			present invention are one or more	AI751250,	AI310126, AI188971, AA906201,
			polynucleotides comprising a	AA019739,	AW264561, AW009062, AI361312,
			nucleotide sequence described by	AA887119,	AA971980, AI580662, AA088862,
			the general formula of a-b, where a	AI261311,	AA575958, AA018414, AI268976,
			is any integer between 1 to 1785 of	AA904689,	AI784506, AI654089, AA838000,
			SEQ ID NO:1909, b is an integer of	AI800634,	AA018103, AA833673, AA809439,
			15 to 1799, where both a and b	AA970480,	AI419770, AW189948, AI806808, N40196,
			correspond to the positions of	AA886637,	H38658, AA059058, AA809455, AA532665,
			nucleotide residues shown in SEQ ID	AI538082,	AA887381, T50287, AI083552, T47520,
			NO:1909, and where b is greater	AA054140,	AA469072, AI933491
			than or equal to a + 14.	AA634291,	N58823, AI799084, H86061, R24685,

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				AI807280,		Z21231, AA019783, H78462,	· Æ	
				AI338489,		W01156, AA016261,	N28787,	
				AF151877,		AL117550,	ഗ	
1910	HTELO87	877881	7	AA115605,	AIS89156,	AA115471,	AI359615,	
			present invention are one or more	AA115213,	AI817096,	N50090, AW118065,	W118065, AI024233,	
			polynucleotides comprising a	AA423826,	AA610042,	A1672797,	AA307285,	-
			nucleotide sequence described by	AI800760,	AA989046,	ř		
			the general formula of a-b, where a	AW162429,	N50523, AA034218,		AA805237, AA115129,	
				AA721969,	AA496544,		AA419084, AA708005,	
			SEQ ID NO:1910, b is an integer of	AI741973,	AI204382,	AA476516,	R70914, R70913,	
			15 to 1267, where both a and b	AA043558,	AA320866,	AA476416,	AA033534,	
				AA781036,	AI627278,	AA903019,	AA347354,	
			nucleotide residues shown in SEQ ID	AA035548,	D25909, AZ	1043557, A.	D25909, AA043557, AI419107, AI080319,	
			NO:1910, and where b is greater	H97516, C	C21455, N50579,	579, AW299!	AW299563, AA310893,	
			than or equal to a + 14.		AI761872,	AA035038,	AA905739,	
				AA746181,	AI521292,	AI554821,	AI433157,	
				AI889189,	AI866469,	AI815232,	AW086285,	
				AI927233,	AI366900,	AI539707,	AI355779,	
				AI590043,	AI440239,	AI537677,	AI494201,	
				AI500659,	AI539800,	AI866465,	AI801325,	
				AI500523,	AI538850,	AI702065,	AI582932,	
				AI923989,	AI872423,	AI284517,	AIS00706,	
				AI491776,	AI445237,	AW151138,	AI521560,	
				AI500662,	AW172723,	AI284509,	AI440263,	
				AI538885,	AI889168,	AI866573,	AI828574,	
				AI633493,	AI434256,	AI434242,	AI805769,	
				AI888661,	AI648454,	AI284513,	AI888118,	
				AI859991,	AI436429,	AI887775,	AI889147,	
				AI581033,	AI371228,	AI567702,	AI440252,	_
				AI866786,	AI610557,	AI860003,	AI242736,	
				AI887499,	AI539781,	AIS00714,	AI559957,	
				AI491710,	AIS21571,	AI582912,	AI623736,	
				AW089557,	AW151974,	AW151979,	AI612913,	
				AI885949,	AI371265,	AL045500,	AI469775,	

AL039390, AI567953, AI446495, AI863014, AI671642, AI890907, AI866581, AI889148, AI285439, AI431307, AI539771, AI804505, AI554827, AI866461, AI815150, AI273179, AI371251, AI866510, AI285419, AI923046, AL047422, AW151136, AI866691, AI924051, AA715307, AI432644, AA809974, AI828583, AI66439, AI872315, AI624545, AI042365,	, AI6485, , AL1336, , AL1336AL133655 , AL133655 , AL1336	I19505, U96138, AL122103, E073 E12888, AL133084, AL133070, AF , AF061836, M30514, Y07915, AR), I96214	AI815614, AA159571, AA , AA485201, H27837, AA , AA384878, W95754, H1	AI802901, AI889514, AA464368, AW026514, AI278645, AA315349, AA777364, AI741517, AW139143, N93194, AA632076, AA700910, AA456473, AI889524, AI160031, AA464386, AA464702, AI089651, AI057409, AI271327, AI921322,
			Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 540 of SEQ ID NO:1911, b is an integer of 15 to 554, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1911, and where b is greater than or equal to a + 14.	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a
			878199	878207
			HWLQL72	нвлгоэ
			1911	1912

			is any integer between 1 to 1704 of	AA417376, AI689262, AA081418, AI611368, R83304,
·			SEQ ID NO:1912, b is an integer of	281824, AA680361, AI
			15 to 1718, where both a and b	AW022859, AW268970, AI273221, AW264836,
_			correspond to the positions of	AW022729, AI184566, AA416981, AW020287, R52791,
			e residue	AI247775, AI924151, AI669435, AI093813,
			NO:1912, and where b is greater	AI206016, AA888936, AW027977, AI269409,
			than or equal to a + 14.	AW027941, AW250197, AI334129, AI474405, N34475,
				AA351606, AA435915, AI270365, AW022849,
				AA650241, AA629813, AA594133, AI358262,
				AA972239, N63595, AI538989, AI075918, AI431608,
	-			AW005549, AI088724, AI240714, AI421046,
	-			
	_			, T06003,
	-			AI767408, AA417194, AA493371, AI688358,
				AW167434, AI688521, AI961941, AW269290,
		_		AA351839, AA024843, AA319841, AA675922, N57835,
				3,
	_			AI566133, AA527515, AA478734, AI700650,
				8, AI393134,
				AA352936, AA364692, AW167540, F09704, AI432014,
		-		AI248967, T66281, AA516011, AI919046, T98208,
				, AA747622, AI523723, AI3485
		-		
	_			0, N42879, AI032060,
_		-		AA642247, AI554380, AW302197,
				AI766194, AW207784, AW376043, C02058, AI033452,
				AC000378, AB019038, Z66003, Z66002, Z65575
HE 1913 HE	HE2HC14 8	878238	Preferably excluded from the	AI127452, AW351965, AW351958, AW178075,
			present invention are one or more	AW351966, AW351967, AW351961, AW177978,
			polynucleotides comprising a	AI659805, AW351960, AA772145, AI336994,
	_	_	nucleotide sequence described by	AW178080, AI332356, AW340996, AW177836,
			al formula of a-b,	AW178082, AW178086, AI703194, AW178079,
			teger between 1 to 1961	AW177841, AA102622, AW136469, AI476336,

	SEQ ID NO:1913, b is an integer of	AI636042,	AW375181,	AW365198,	AI813938,
	to 1975, where both a	AI769135,	AI074596,	AA418593,	AW178083,
-	to the position	AI498407,	AI654773,	AW351962,	AW177876,
	nucleotide residues shown in SEQ ID	AI366827,	AW178077,	AW020441,	AA806382,
	NO:1913, and where b is greater	AW178182,	AW178076,	AW178081,	AW177879,
	equal to a + 14.	AW365184,	AW366023,	AW365168,	AW375184,
		AA418655,	AW177839,	AW178084,	AI468009,
		AI433820,	AI692309,	AW082896,	AI927777,
		AW365192,	AW387262,	AI143953,	AW365194,
		AA421501,	AI271676,	AA425855,	AA854439,
		AW082902,	AW177842,	AW128928,	AI392856,
		AW365398,	AA421470,	AW365185,	AA535678,
		AI400413,	AW365353,	AW387278,	AA680114,
		AI076707,	AI285336,	AW365392,	AI581008,
		AW375185,	AA938196,	AI801859,	AW089786,
		AI382040,	AW365381,	AW365201,	AW375183,
		AI243492,	AA973630,	AL120271,	AA649053,
		AW365405,	AI698558,	AA934487,	AW366025, R98908
		AI473267,	H70023, AA	A976681, AW365408,	W365408, AA80662
		AW375120,	AI536915,	AW178078,	AW365180,
		AW365183,	AW003830,	AW178085	AA400106,
•		AA532939,			W85961, AW387263,
		H58724, A	AI301165, A		AA463549, AA52734
		AW262369,	AI830518,	AA832369,	AI383837,
		AI216813,	AA280430,	AW177877,	AW365189,
		AW177079,	AI288375,	AW375133,	AA515868,
		AW375160,	AW243710,	AW375442,	R98681, AA932395
		AW169226,	AA188895,	AI335817,	AW365411,
		AW365146,	AW365417,	AW382189,	AW365202,
		AW382124,	W24191, A		AI868465, AA280348
		AW365182,	R97677, A	AW365412, H	H56644, W72745,
		AW177846,	AW365404,	AW365404, AW365402,	AW365359,
		AA424055,	AW177974,	AW177974, AW365164,	N91771, AW36519
		AW351813,		120462, AW3	
		AW375130,	R84876, A	AW365362, C	C01884, AW351560,
		AW375422,	AW365364,	AW366058,	AA936703, AC008040

α	878774	Droforably ovaluded from the	
))		invention are on	
		leotides comprising a	A1034324, N80327, AA350189, AW351942, AA349355, W04179, AF201978, H144879, H43607, H43649
		nucleotide sequence described by	
-		the general formula of a-b, where a	
		SEQ ID NO:1914, b is an integer of	
		15 to 508, where both a and b	-
		correspond to the positions of	
		·O	
		-	
	_4	than or equal to a + 14.	
878374			AL041566, AA477266, AI656936, AI951716,
	_	present invention are one or more	AI096374, AA477267, AI927648, AA292231.
	_	polynucleotides comprising a	
_		nucleotide sequence described by	AI15039
	-	the general formula of a-b, where a	AA42752
·	•-	teger between	AA293470, AA723836, AA99409
	נט	SEQ ID NO:1915, b is an integer of	1985377, H4
_	_	15 to 2885, where both a and b	
0	U	correspond to the positions of	R06788, AA808474, T79352, Z45236, F04128,
<u> </u>	드	de residues a	AA503842, AI361214
Z -	z .	, and w	A987751, R00061,
	יי	than or equal to a + 14.	, AI424488, F08632, AA293015
			AA693978,
		-	R06739, AA343968,
\dashv	- ['		AA226870, AB033010, AL137675
878403		፫ .	AL048840, AI064902, AW249691, AI872413,
	_	present invention are one or more	
		polynucleotides comprising a	AW004004, AI923006, AA587051, AA279533,
		nucleotide sequence described by	AW183520, AI419833, AW292319, AA214039,
		the general formula of a-b, where a	, AW16706
	• •	teger between 1 to 2994	
		SEQ ID NO:1916, b is an integer of	AW088356, AI336423, AI803586, AA100821,
	٠.١	15 to 3008, where both a and b	AL048839, AW105007, AA332665, AW021472, W93478.

			correspond to the positions of nucleotide residues shown in SEQ ID	AA211303, R51407, AA040271, AI128507, AI824743, AI520729, AA279532, N62195, AA770032, AI991817,
			nd where b	W67473, AA309583, AW392599, AA976795, R14643,
			than or equal to a + 14.	AA9/6594, AL216/60, AA4429/2, K5356/, AA36989/, AI364305, T56013, AW021133, AA016204, R53679,
				H73568, AI521207, AA
				AI832743, AA609475,
	-			AA563648, AI824485, AI561042,
				AA040252, AI383108, AA579428, AA305720, T91394, T04986 D45624 T86544 D29736 C00010 T29665
				T05066, AA887773, AI985106, T85482, AW243484,
				AA720874, AA573214,
				AA305679, L25798, X66435, AL079334, AL050004,
1917	HTPAY82	878433	Preferably excluded from the	
			present invention are one or more	AI128804, AI826623, AA516431, AI989747,
			polynucleotides comprising a	AW183193, AI141284, AI989739, AA702011,
			nucleotide sequence described by	AA911088, AA989129, AA876539, AA477156,
			the general formula of a-b, where a	AA305052, W19506, N89912, AI265924, AA644621,
			is any integer between 1 to 544 of	2820, AI633679, AA987264,
			SEQ ID NO:1917, b is an integer of	
			15 to 558, where both a and b	W93906, AI198595, AA946978, AI419292, AI198127,
			correspond to the positions of	
				AA927461, T97984, AA341602, AA035640, AA356704,
			NO:1917, and where b is greater	AI933253,
			than or equal to a + 14.	AA780176
				AA338761, AA234074, T98061, T83106, AA193255,
				AF096895, AF057306, AF135380, AF135381, AF145216
1918	HMUBQ39	878436	Preferably excluded from the	AW084650, AA088424, AI697069, AA172042,
			present invention are one or more	, AA172044, AI744623,
			polynucleotides comprising a	, AA993207, AI371167,
			nucleotide sequence described by	AI890821, AA609797, AI018761, AW372890,
			the general formula of a-b, where a	AI814927, AA625264, AI954856, AA993191,
			is any integer between 1 to 1805 of	AA614086, H05584, AI961696, R39132, AI632376,
			SEQ ID NO:1918, b is an integer of	AI143462, AW136636, AA722935, AA172197, D20763,

			15 to 1819, where both a and b	AA701379, F06989, AA148617, AW044004, R21296,
			correspond to the positions of	AA172201,
			de residue	AI627401, N42449, AI224491, AA635934, R14008,
			NO:1918, and where b is greater	H05119, R18980, T26664, T16725, F07496, T59139,
			than or equal to a + 14.	AA372447, AA092086, F31653, Z40099, AW271655,
				AA993655, R32993, R46141, AI472512, T59062,
				T26665, Z40560, R32717, AA148756, AA374317,
				AA585413, AA064920, AI917682, AA625242, R32994,
				AW362703, AW372891, AW386147, R25109, R25628,
	-			AA828475, R3
				AIS40458, AI814841, AIS70152, AW079699,
	,			85, AA836253,
				AI954475, AI689096, H03560, AI368579, AI357049,
				AI560184, AI469505, AI687295, AA767252,
				AI280732,
				AA923096, AI341690, AI888575, AI697178,
	-			AI765469, AW075921, R30844, AI702494, AI359787,
				AI417754, AW104141, AI867017, AA742592,
				9, AA741502,
				AI679261,
				7, AI749231,
				AI623980, AI590755, AC0052
				AF102578, AF038847, U67810, A85213, AB015752,
				AL137490,
				AF039907, AL049552
1919	HCEYN60	878560	Preferably excluded from the	
			present invention are one or more), T46897, R40801
			polynucleotides comprising a	7
			nucleotide sequence described by	H49579, H49658,
			the general formula of a-b, where a	H62359, N23682, AA002170, AA039225, AA045879,
			ger between 1 to 563	AA0534
			SEQ ID NO:1919, b is an integer of	AA171927, AA173260, AA181967, AA186968,
			15 to 577, where both a and b	AA215430, AA215576, AA494375, AA554350,
			correspond to the positions of	AA565187, AA582635, AA594327, AA612625,
			nucleotide residues shown in SEQ ID	AA878313, AA886926, AA887637, AA908475,
			NO:1919, and where b is greater	AA939096, AI051140, AI083860, AA641276,

			than or equal to a + 14.	AA205608,	AA284538, AA411196, AA410243,
				AA411096,	AA436335, AA478263, AA478319,
				AA609270,	AA628990, Z19827, AA719345, AA769770,
				AA776741,	AI018379, D19640, AI305530, AI307824,
	•			AI344950,	AI349732, AI363496, AI368551,
	-			AI434470,	AIS61271, AI498585, AI423077,
				AI147393,	AI167340, AI224833, AI174303,
				AI187983,	AI659839
1920	HWHGF46	878800	Preferably excluded from the	AA814195,	AI457718, AI085388, AI765650,
			present invention are one or more	AA633558,	AI379449, AI476182, AI419034,
			polynucleotides comprising a	AI037888,	AI148797, AA028963, AW009541,
			nucleotide sequence described by	AW051402,	
			the general formula of a-b, where a	W67782, AA035136,	A035136, AI016426, AI304821, AA085457,
			is any integer between 1 to 2101 of	AI808210,	AA098932, AI685969, W39585, AI685970,
			SEQ ID NO:1920, b is an integer of	AI038819,	AI219571, AI580447, AA485877,
			15 to 2115, where both a and b	AA487780,	W42434, AA594455, AI865081, AI085147,
			correspond to the positions of	AI202241,	AA632996, AA035135, D45612, AA991990,
			nucleotide residues shown in SEQ ID	AC006261,	AL031985, AL021154, AC006449,
	•		NO:1920, and where b is greater	AL008718,	Z95329, AC004950, AC002349, AL031846,
				AF146367	
1921	HPMSF50	878909	Preferably excluded from the	AL045860,	N58437, AI525782, AI688578, AA007479,
			present invention are one or more	AA310929,	AA906018, N41678, AW084721, N59420,
			polynucleotides comprising a	AA007400,	AA234496, AI810048, AI394367,
			nucleotide sequence described by	AW273848,	AI400139, AI659487, AI168584,
			the general formula of a-b, where a	AW247506,	•
			is any integer between 1 to 3939 of	AA235036,	
			SEQ ID NO:1921, b is an integer of	AA630558,	
			15 to 3953, where both a and b	AA877580,	AA931472, AA351722, AA232945,
	-		correspond to the positions of	AI208004,	AA885392, N71533, H09450, AA554688,
			nucleotide residues shown in SEQ ID	AA983994,	AI221004, AA235204, H54147, AA460203,
			NO:1921, and where b is greater	AA985683,	AI681824, N22166, AA889639, AA668373,
			than or equal to a + 14.	H81138, AA678603,	A678603, R97728, AW291709, AI346634,
				AA337087,	T56721, C14300, AA310347, AA359522,
	•			AI032752,	AI032752, AA705700, R68352, R10225, C14263,
	-			T40018, H	T40018, H81043, T56722, C14304, R68562,
				AI369399,	R96796, AA333514, AA459932, H57429,

			AI758833, AA836349, C14291, AA902529, C14302,
			9495, R10732, N937
-		-	AA665646, R12861, AA384438, AA682859, AI904934,
			4935, D80004
HTWEA61	878917	Preferably excluded from the	
		present invention are one or more	3,
		polynucleotides comprising a	AW192514, AI566340, AI972077, AI811155,
		nucleotide sequence described by	A1936746, A1089502, AI372947, AI004230,
		the general formula of a-b, where a	AI354532, AL119666, AI084362, AI027083,
		is any integer between 1 to 1978 of	AI691080, AA621070, AI744332, AI149953,
		SEQ ID NO:1922, b is an integer of	
		15 to 1992, where both a and b	AI003733, W20002, AW074007, AI627187, AW242075,
		correspond to the positions of	AW130451, AI014764, AI091649, AA041468, W55944,
		NO:1922, and where b is greater	AA040575, AI689545, AI524423, AI521587,
		than or equal to a + 14.	_
			AI538583, AW263138,
			AI368864, AW316596, AI539834, AI952557,
			AA662403, AW
			AA808860, N78681, N32970, AA176087, AI125767,
	_		, AI074758
			, AI372493,
. <u>-</u>			
-			œ
	-		D82747, W26208, AA471277, AA903068, AI680414,
			AL038664, AA664940, AA897635, AI535982, D31438,
			AI419708, AW275741, AA386197, R62151, AI051237,
			R62259, W28043, R39290, AI250661, F10830,
			AI695489, AA343846, R43842, AA334321, AA093703,
			AA845417, AA332748,
			AI524545, AA095572,
			24108, AI611841, AA
			AA323934,
			AW021288, AA329440, D81428, AA344329, AA039822,
		-	AW375337. AW270647. AW149580. F35697. AM148718

				AA862706, AI523217, T25879, R1 T69962, T7 AA093662, AA349447, AA199620, U85195, AE		AA848160, AI241878, '0902, AA71 '6163, T699 '0652, AW27 AA373966, AIS57186,	
1923	HILBF77	878931	preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 711 of SEQ ID NO:1923, b is an integer of 15 to 725, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1923, and where b is greater than or equal to a + 14.	AW242021, AL049923	AA352298, AA330358,	AA330358,	Z78381, C01470,
1924	нтенхоз	879009	Δ 4 4 0 4 Z N O 4 .	AI872206, AW004890, AA831357, AA581345, AI360561, AI277190, AI522238, AA706811, AW450726, AI702026, AI624976, AF035606,	AI912340, AI572080, AW074361, AI690445, AW439592, AA100279, AI015234, AI744762, AA122332, AA122332, AI681670, US8773	AI758821, AW058001, AI361820, AI917776, AI798286, AA485257, AH689240, AW265061, T34498, AI AA092467, AA092467,	AI758821, AW337178, AW058001, AA75261, AI361820, D20022, AI982775, AI91776, AA825538, AI798286, AI140796, AA485257, AA835492, AA689240, AI469550, AW265061, AI884872, T34498, AI811224, AI355770, AA092467, AI597962, AA089786, AA654171,
1925	HPHAA47	879234	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by	AIS40230, AI346427, AW131500, AI089921,	AI453545, AI819403, AI419533, AI347957,	AI697681, AI857677, AW027758, AA612573,	AW170551, AI348016, AW016071, AI601101,

		the general formula of a-b, where a	AI088798, AI123932, AI348513, AA916423,
		3897 0	AI697840, AI346773, AI82727
		ы	AI763320, AA609447, AA02442
		15 to 3911, where both a and b	AA948406, AW149724, AI435604, AA946618,
		correspond to the positions of	
		ס	AI814488, AA232203, H43798, AW374530, F11803,
		and where b	W131707, AI285224,
		than or equal to a + 14.	H23601, N51357, AA
			AA758706, AI927091, Z39461, AA936791, H23640,
			H43806, AA364902, AI802791, AA864755, T33777,
	-		F02788, H42258, F09452, AA583801, T65604,
			R43369, T65538, H40427, AA336254, W94547,
_			
			AA580399, W78003, AA463368, AW293983, AW374487,
			AA513346, N29649, AA837760, AA024429, AI695172,
			R17652, AW448962, AA232743, AA973192, AA652557,
-			W79462,
			R65673, AA719939, X85665, AI972788, AI972806,
			AA933622, AA916725, AW006745, AL137343
1926 HHFJJ61	879386	Preferably excluded from the	R93802, AA130402, H07960, AW250644, H85944,
-		present invention are one or more	R85969, AA095215, AA036855, AA215398, AA308813,
_		polynucleotides comprising a	1, AA324032
		nucleotide sequence described by	
		the general formula of a-b, where a	
-		is any integer between 1 to 1027 of	
		SEQ ID NO:1926, b is an integer of	
		15 to 1041, where both a and b	
		correspond to the positions of	
		nucleotide residues shown in SEQ ID	
		NO:1926, and where b is greater	
		than or equal to a + 14.	
1927 H2CAA49	9 879484	Preferably excluded from the	AI279876, AI539769, AA876127, AI963800,
		present invention are one or more	AA206425, AI969470, AI951966, AA459503,
		polynucleotides comprising a	AA778294, AA639198, AA446426, AI334209,
		nucleotide sequence described by	AI150191, AI281280, AW149760, AA446118,

is any integer between 1 to 2296 of AN376 SEQ ID NO:1927, b is an integer of AN376 Is to 2310, where both a and b Correspond to the positions of C7560 nuclectide residues shown in SEQ ID NO:1927, and where b is greater T6553 than or equal to a + 14. T6553 AN396 AA503 AA503 AA503 AA504 AA506	is an integer of AW376909, AI127770, AI139373, AI753243 both a and b AA789258, N95643, AI754062, AA236574, positions of AL121103, AA213367, AA837311, AI187231 AA27539, AI344110, H67810, W95535, AI T65536, AA872668, AI192986, CI7463, AI AA770471, T17222, AW192135, AA075621, AA470471, T17222, AW192135, AA075621, AW139044, AI1913866, AA192466, AA165156 AI826398, AA678954, AI271344, AA113939 AA137249, H17790, F11801, AA164768, C7 R89384, T16445, T69722, N66040, C18698 AA503343, AA339152, AI625443, D81644, H58956, D60375, F06655, H58600, AA5146 H02142, AA164700, AA055768, AA306967, AI568159, C21496, W95420, H68082, AI57841, AA536620, AW363691, AA142866, R01641, AA524392, T85647, Z39669, H17791, H586
is an integer of both a and b positions of is shown in SEQ ID is b is greater i + 14.	is an integer of AW376909 both a and b AA789258 positions of C75603, s shown in SEQ ID AL121103 AA27535; t + 14. AA57836, AA58984, AA58626, H02142, AA56159
both a and b positions of s shown in SEQ ID b is greater + 14.	both a and b positions of s shown in SEQ ID bis greater t + 14. AA470471 AW139044 AR82538 AA137249 R89384, AA503343 H02142, AA503343 AA58256, AA588562
positions of shown in SEQ ID be is greater to the second t	positions of C75603, s shown in SEQ ID AL121103; b is greater T6536, 1 + 14. T6536, AA470471 AW139044 AR137249 R89384, AA503343 H02142, AA5033626
s shown in SEQ ID b is greater t + 14.	s shown in SEQ ID AL121103 b is greater AA227536, 1 + 14. T65536, AA470471 AW139044 AN39044, AN37245 AA37245 AA503342, AA503342, AA503342, AA503342, AA503342, AA503342, AA382754
and where b is greater equal to a + 14.	greater AA227536, AA470471 AW139044 AN826398 AA137245 R89384, AA503347 H58956, H02142, A1568155 AA382755
equal to a + 14.	A470471 AW13904 AW13904 AN13904 AN13904 AN137249 R89384, AA503343 H58956, H02142, AI568159 AA382754 AA524392
AA470 AW139 AV139 AV130 AV137 R8938 AV20 AV214 AV20 AV20 AV20 AV20 AV20 AV20 AV20 AV20	AA470471, T17222, AW192135, AA075621, AA506763, AW139044, A1913866, AA192466, AA165156, AA1826398, AA678954, AI271344, AA113939, C05669, AA137249, H17790, F11801, AA164768, C75565, R89384, T16445, T69722, N66040, C18698, H59003, AA503343, AA339152, A1025443, D81644, R78076, H58956, D60375, F06655, H58600, AA514607, H02142, AA164700, AA055768, AA306967, T70379, A1568159, C21496, W95420, H68082, A1572235, AA38620, AA36620, AA36620, AA36620, AA36620, AA36620, AA36620, AA36691, AA142866, R01641, F09450, AA524392, T85647, Z39669, H17791, H58601,
AM139 A1826 AA137 R8938 AA503 H5895 H5895 H7885 AA236 AA382	AW139044, AI913866, AA192466, AA165156, AI826398, AA678954, AI271344, AA113939, C05669, AA137249, H17790, F11801, AA164768, C75565, R89384, T16445, T69722, N66040, C18698, H59003, AA503343, AA339152, AI025443, D81644, R78076, H58956, D60375, F06655, H58600, AA514607, H02142, AA164700, AA055768, AA306967, T70379, AI568159, C21496, W95420, H68082, AI572235, AA382754, AA989472, T35523, H02038, T65602, AA236620, AW363691, AA142866, R01641, F09450, AA524392, T85647, Z39669, H17791, H58601,
A1826 AA137 R8938. R8938. AA503 H5895 H0214 A1568 AA382 AA382 AA382 AA458	AI826398, AA678954, AI271344, AA113939, CO5669, AA137249, H17790, F11801, AA164768, C75565, R89384, T16445, T69722, N66040, C18698, H59003, AA503343, AA339152, AI025443, D81644, R78076, H58956, D60375, F06655, H58600, AA514607, H02142, AA164700, AA055768, AA306967, T70379, AI568159, C21496, W95420, H68082, AI572235, AA382754, AA989472, T35523, H02038, T65602, AA35620, AW363691, AA142866, R01641, F09450, AA524392, T85647, Z39669, H17791, H58601,
AA1377 R8938. AA503 H5895 H0214 AA185 AA236 AA236 AA382 AA165 AA165 AA165 AA165 AA165 AA165 AA165 AA165 AA165 AA1695 AA1933	AA137249, H17790, F11801, AA164768, C75565, R89384, T16445, T69722, N66040, C18698, H59003, AA503343, AA339152, AI025443, D81644, R78076, H58956, D60375, F06655, H58600, AA514607, H02142, AA164700, AA055768, AA306967, T70379, AI568159, C21496, W95420, H68082, AI572235, AA382754, AA989472, T35523, H02038, T65602, AA236620, AW363691, AA142866, R01641, F09450, AA524392, T85647, Z39669, H17791, H58601,
R8938 AA503 H5895 H0214 A1568 AA382 AA455 AA455 AA465 AA65 AA	R89384, T16445, T69722, N66040, C18698, H59003, AA503343, AA339152, AI025443, D81644, R78076, H58956, D60375, F06655, H58600, AA514607, H02142, AA164700, AA055768, AA306967, T70379, AI568159, C21496, W95420, H68082, AI572235, AA382754, AA989472, T35523, H02038, T65602, AA36620, AW363691, AA142866, R01641, F09450, AA524392, T85647, Z39669, H17791, H58601,
AA503 H5895 H0214 A1568 AA382 AA524 AA524 AA65 AA165 AA029 AA165 AA165 AA165 AA165 AA165 AA165 AA165 AA1695 AA1933	AA503343, AA339152, AI025443, D81644, R78076, H58956, D60375, F06655, H58600, AA514607, H02142, AA164700, AA055768, AA306967, T70379, A1568159, C21496, W95420, H68082, A1572235, AA382754, AA989472, T35523, H02038, T65602, AA356620, AW363691, AA142866, R01641, F09450, AA524392, T85647, Z39669, H17791, H58601,
H5895 H0214 A1568 AA382 AA236 AA524 AA382 AA165 AW029 T8210 Z2108 H7748 H7748 AW385 AW8080 AI695 AI693	W K
H0214 A1568 AA382 AA236 AA236 AA382 AA165 AW029 T8210 Z2108 T9726 AA385 AW385 AW385 AW385 AW385 AW385 AW385 AW385 AW385 AN593	A164700, AA055768, AA306967, C21496, W95420, H68082, AI57 AA989472, T35523, H02038, T6 AW363691, AA142866, R01641, T85647, Z39669, H17791, H586
A1568 AA382 AA236 AA524 AA524 AA524 AA165 AA165 AA165 AA165 AA165 AA165 AA165 AA1695 AA1695 AI1695	C21496, W95420, H68082, AI57 AA989472, T35523, H02038, T6 AW363691, AA142866, R01641, T85647, Z39669, H17791, H586
AA382 AA524 AA524 AA382 AA165 AA029 TR210 Z2108 T9726 T9726 AA385 AW385 AW385 AW385 AW1695 AI695	AA989472, T35523, H02038, T6 AW363691, AA142866, R01641, T85647, Z39669, H17791, H586
AA236 AA524 AA382 AA165 AW029 TR210 Z2108 H7748 H7748 T9726 AA934 AW385 AW080 AI695 AI693	AW363691, AA142866, R01641, T85647, Z39669, H17791, H586
AA524 AA382 AA165 AW029 T8210 Z2108 H7748 H7748 T9726 AA934 AW385 AW080 AI695 AI693	T85647, Z39669, H17791, H586
AA382 AA165 AW029 T8210 Z2108 H7748 T9726 AA934 AW385 AW080 AI695 AI695 AI693	
AA165 AW029 T8210 Z2108 H7748 H7748 T9726 AA934 AW385 AW080 AI695 AI600 AI693	
AW029 T8210 Z2108 H7748 H7748 T9726 AA934 AW385 AW385 AW1895 AI695 AI693	AA165228, AA838767, AA165229, R42323, AI025112,
T8210 Z2108 H7748 H7748 T9726 AA934 AW385 AW385 AW1895 AI695 AI693	AW029182, AA865982, T91320, C00668, T99684,
22108 H7748 T9726 AA934 AW385 AW080 AI695 AI693	T39127,
H7748 T9726 AA934 AW385 AW080 AI695 AI695 AI693	
T9726 AA934 AW385 AW080 AI695 AI695 AI6933	R16380, AA937248, AA199583, AA
AA934 AW385 AW380 AW080 AL695 AI400 AI5933	
AW385 AW080 AI695 AI400 AI933	AA327356, T87388,
AW080 AI695 AI400 AI933	AW385433, AW385409, Z20096, AI924498, AA513297,
A1695 A1400 A1933	AW080588, AA558986, AI926128, AI581525,
A1400	AI695291, AW196067, AI783818, AI623264,
	AI400863, AA526975, AI445127, AI469613,
	AI933636, AI919084, AA632103, AA581848,
AI888	AI358508, AI469656,
AI275	5, AI249798, AA552670,
A1040	AI040152, AI242802, AA884931, AI378681,

				AI025266,	AI434099,	AA533047,	AW272720,
		_		AI801054,	AI888914,	AI735767,	AW304001,
				AI445913,	AI436796,	AW190856,	AI921153,
				AI380637,	AI888294,	AI634717,	AI075324,
				AI815198,	AI805627,	AI932444,	AW073291,
				AI891014,	AA425142,	AA622524,	H67122, AI916480,
				AI146786,	AA316874,	AI678847,	AA315049,
				AI817063,	AA573742,	AW152548,	AW151674,
		_		AI610106,	AI675865,	AW152169,	AI675714,
				AW027843,	AI475938,	AI685830,	AA582017,
				AI473626,	AW381550,	AI445130,	AI800451,
				AI800431,	AI972701,	AI678427,	AI801784,
		_		AI582452,	AI867585,	AI972499,	AI720013,
				AI278406,	AI277266,	AI082505,	AW191880,
				AI537173,	AI473553,	AI925030,	AI559391,
				AI471336,	AF053641,	U33286, AI	AF038452, AF053642
				AF053650,	AF053651,	AF038451,	AF053640,
				AF007791,	AF088867,	AA570120	
1928	HCRNW08	879595	Preferably excluded from the	AA192153			
			present invention are one or more				
_			polynucleotides comprising a				
			nucleotide sequence described by				
			the general formula of a-b, where a				
			is any integer between 1 to 407 of				
			SEQ ID NO:1928, b is an integer of				
			15 to 421, where both a and b				
			correspond to the positions of				
			nucleotide residues shown in SEQ ID				
			NO:1928, and where b is greater				
			than or equal to a + 14.				
1929	HNTD129	879661	Preferably excluded from the	AA555115,	AW083142,	AW383992,	AI819977,
			present invention are one or more	AI818981,	AW302146,	AI357211,	AA970333,
			polynucleotides comprising a	AA565308,	AW391496,	AA809752,	AA043134, C18608
			nucleotide sequence described by	AA548230,	AA565317,	AI352620,	AA554155,
			the general formula of a-b, where a	AA279358,	AW392424,	AA043611,	AI433904,
			er between	AA767874,	AA370804,	F33509, A	AW370978, AI500136

			SEQ ID NO:1929, b is an integer of	AA360902,	AA279306,	AA370803,	AA360902, AA279306, AA370803, AC004677, AL078630
			both a and				
			correspond to the positions of				
			nucleotide residues shown in SEQ ID				
			e p,				
			than or equal to a + 14.		- 1		
1930	HCRNM29	879886	Preferably excluded from the	AA040621,	R64534, A	AA811265, AI	
			present invention are one or more	AI222332,	AA040620,	AA040620, AW001618, N40203,	N40203, AI796277
-			polynucleotides comprising a				
			nucleotide sequence described by				
•	-		the general formula of a-b, where a				
			is any integer between 1 to 748 of				
			SEQ ID NO:1930, b is an integer of				
_			15 to 762, where both a and b				
			correspond to the positions of				
			nucleotide residues shown in SEQ ID				
			NO:1930, and where b is greater				
			equal to a + 14.				
1931	HTPAM76	880071		AW387764,	AW387814,	AW387802,	AW387787,
			present invention are one or more	AW387847,	AI888586,	AW387804,	AA156240,
			polynucleotides comprising a	AA156243,	AA115637,	AW388637,	AW387768,
			nucleotide sequence described by	AW073692,	AW387860,	AI828610,	AA447697,
			the general formula of a-b, where a	AW078652,	AA156747,	AW387867,	AA115638,
			is any integer between 1 to 1619 of	AW387851,	AA147510,	AW387845,	AA147381,
	_		SEQ ID NO:1931, b is an integer of	AI671236,	AA627367,	AI302358,	AW387765,
			15 to 1633, where both a and b	AI589344,	AA126967,	AW194339,	AA552339,
			correspond to the positions of	AW274844,	AA115437,	AA631614,	AA482223,
			e residue	AI336522,	AI610638,	AA464766,	AA127119,
			NO:1931, and where b is greater	AA148915,	AI801445,	AI888444,	AA486631,
			than or equal to a + 14.	AA481927,	AI926413,	AW058286,	AA468787,
				AA156919,	AI888332,	AA115436,	AW387859,
				AA129137,	AA911832,	AA480064,	AW387887,
				AI446210,	AA129136,	A1935846,	T93584, AW338675,
				AA486537,	AA447849,	AA373191,	AI739001,
				AI536744,	AA300698,	AI926870,	T79051, AW378720,
				T70156, AV	AW387878, A	AW150592, A	AI805203, AI678275,

				אנאנאנין אפאנוט אסטטען דאסטטן אסטטטטטטטטטטטטטטטטטטטטטטטטטטטטטטטט
				AA649486, AA652093,
_	-			5, AA364232, AI654194,
				AA143334, AA372265, AA026564, N78458, AI472423,
				AA026472, AA313840, N55383, AF112214, D17244,
				D17071, AA706862
1933	HLSAA96	880418	Preferably excluded from the	AW444874, AI920970,
			present invention are one or more	AA431746, AA651708, AA847822, AA746501,
			polynucleotides comprising a	AI051249, AI005487, AI368709, AI417856,
			nucleotide sequence described by	AA009824, H06206, AW150601, H08319, AA830175,
			the general formula of a-b, where a	AA809393, AA765426, AW337780, AI435979,
			is any integer between 1 to 1783 of	, AA508643, AA2826
				R44287, R59778, AA768684, AI193720, AW235814,
			15 to 1797, where both a and b	AA993048, R61320, T09292, AA503026, AA301325,
			correspond to the positions of	AW084853, H08221, T84812, T78009, AA340198,
			nucleotide residues shown in SEQ ID	AA009714, R23537, AI933451, AA649008, AA322332,
			NO:1933, and where b is greater	
			equal to a + 14.	
1934	HBBMA61	880578	bly exc	AA934705, AI370920, AI744886, W86237, AA609163,
			present invention are one or more	AI082256, AI140436, N53361, AA968467, AI216727,
			polynucleotides comprising a	N62199, AI143325, AI015198, AW236133, AA732867,
			nucleotide sequence described by	AW341974, AI591092, AI141509, AA002163, N36129,
			the general formula of a-b, where a	R45071, R07479, Z38172, AA059224, T33713,
			is any integer between 1 to 323 of	AI469204, D11576, D11575, Z78385, N64142,
			SEQ ID NO:1934, b is an integer of	T31044, AW243169, AA844013, AA417247, AL119457,
			15 to 337, where both a and b), AL119324
			correspond to the positions of	U46350, AL119483, AL119319, U46347, AL119399,
			nucleotide residues shown in SEQ ID	AL119484, AL119391, AL119418, Z99396, AL134531,
			NO:1934, and where b is greater	AW372827, AW384394, AW363220, AL134533,
			than or equal to a + 14.	AL119363, AL119355, U46349, AL119522, U46341,
				AL119439, AL119444, AL134538, AL119341,
				AL037205, AL119401, U46346, AL119335, AL119396,
				AL119496, AL134920, U46345, AF090190, AB026436,
				AR060234, AR066494, AR054110, A81671, AR069079
1935	HE8QG48	880649	Preferably excluded from the	AA984117, AW163623, AA311680, AA418057,
	,		present invention are one or more	AI144311, AL120308, AA056148, AA187561,

			polynucleotides comprising a	AF072813, W	W01018, AA992009, AA325639, W19986,
			nucleotide sequence described by	AA776635, T.	T30663, T33734, AI878939, AA256403,
			the general formula of a-b, where a		05294, AA134519, Z43583, AA227076,
			teger between 1 to 1310	F06381, AWZ	AW204252, AA430244, AA938909, H30186,
			SEQ ID NO:1935, b is an integer of	D58629, R52	D58629, R52851, N98255, AA161199, AA100159,
			15 to 1330, where both a and b	AA114264, H.	AA114264, H43926, R22746, R34517, AA233577,
			correspond to the positions of	AA081447, A	AA324916, AW138505, AA157365,
			ъ.	AA324268, H	AA324268, H84964, AA019377, AA232373, H42692,
			NO:1935, and where b is greater	W28863, N83	N83234, AA233594, R17978, W81009,
			than or equal to a + 14.	W99386, T34	T34516, T35956, AA214355, AA324917,
				N42109, AAO	
				AA094192, T.	T32010, T31224, Z39649, T87432,
				R22276, AA3	R99404,
				H39131, R16	AA984677
			-	AI755053, A	AI755053, AA362885, AA354497, AA918044, T34825,
				AA417901, A	AA134510, AA643681, AA579642, T34772,
•				AI147468, A	AI336174, AW374188, H19354, AA357382,
				N55823, AA4	N55823, AA482456, AW273035, AA161200, AI911850,
				AA565772, A	AI276668, AA575906, AW337856,
					AA256297, AI308794, AA587048,
				AI354787, R	R99312, AA626391, AF119297, AF059524,
				AR028523, AI	AF059529, AF059525, AF059527,
				AF059526, U	U25265, AF059528
1936	HHENW13	880694	Preferably excluded from the	1,	AI991002, AW087339, AA464410, W37647,
			present invention are one or more		, AA581972,
					AW276040, AI125339, AA167314,
			nucleotide sequence described by		AI803380, AA313202, AI264016,
			the general formula of a-b, where a	-	AW167731, AI083960, AI991293,
			is any integer between 1 to 664 of	AI038896, AV	AW205414, AI460022, AA694199,
			SEQ ID NO:1936, b is an integer of	AI610383, A	AI707649, AI277698, R53610, AA305224,
			15 to 678, where both a and b	AW079550, AJ	AA430117, AA577381, AI074864, N23143,
			correspond to the positions of	AA860618, A	
			nucleotide residues shown in SEQ ID	AI151318, WJ	W16866, R50528, R55254, AA135047,
			NO:1936, and where b is greater	AA255556, A]	
			than or equal to a + 14.	AL040668, W	W37383, AA844913, W93357, R50622,

				N79251, AW271218, AA908394, AI214414, R51941,
				AI669222,
				7, T30416,
				93445, AAO
				AA033670,
				AB001740, AB012865, AB012727
1937	HE8SB64	880747	Preferably excluded from the	AW070902,
			present invention are one or more	AI417256,
			polynucleotides comprising a	AW383890, AI565996, AI499115, AW383902, N21309,
-			nucleotide sequence described by	AA147128, AI767271, AA885289, AI750960,
			a]	AI276772, AW102917, N46066, AI290500, H99543,
			is any integer between 1 to 2414 of	AI302412, AI246663, AL046164, AI242761, N31244,
·				
				M91216, H80005, H85099, AA226631, AI436734,
-			correspond to the positions of	AA460989, D29810
-			nucleotide residues shown in SEQ ID	
			NO:1937, and where b is greater	
			equal to a + 14.	
1938	HKAEN78	880927	Preferably excluded from the	AA306924, T73855, T83294, T85637
			present invention are one or more	
			polynucleotides comprising a	
			nucleotide sequence described by	
			formula of a-b,	
			is any integer between 1 to 908 of	
			SEQ ID NO:1938, b is an integer of	
			15 to 922, where both a and b	
			correspond to the positions of	
			О	
			NO:1938, and where b is greater	
			than or equal to a + 14.	
1939	HOSML44	880994	Preferably excluded from the	AA522719, AA905625,
			present invention are one or more	AI418276, AI560743, AW130435, AI992293,
			polynucleotides comprising a	AI800639, AI204546, AA858118, AA813011,
			nucleotide sequence described by	, AA846821,
			the general formula of a-b, where a	AI362691, AI356940, AI149942, AW008254, N55455,
			is any integer between 1 to 742 of	T79403, AI221349, AA975506, W96084, AW020847,

			SEQ ID NO:1939, b is an integer of	AI240036, AI560812, AI300180, AI089271, H54573,
			where	AA232733, T90553, R94479,
			correspond to the positions of	AW026456, AA768615, AA854918, T
			$\boldsymbol{\sigma}$	W96085, R08289, R94069, H60026, AI685154,
			, and where b i	AW261910
		*	than or equal to a + 14.	, AA883234, N80142,
				A906638, AA995348,
				AI240974, AA738193
				AA443008, N35116, H54683, AW128861, N23206,
				AA364712, AA402136, H96792, AI906874, AI025840,
				AI346239, D59957, H24210, H95663, N20084,
				NZ
				H96607, N90414, T56966, R20754, AA384679,
				AI027068, AI370536, AI520954, T78586, R20753,
				D60276, AI362623, D80608, R54942, AI962075,
				Z28499, H53597, H18631, H91182, H48906,
				3, AA301182, AI985444,
				AA894582, AA609747, AI804799, D59884, AA492083,
				H67369, T27025, H96239
				AA761468, AA972438, AA970691, AA235389,
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				AA319076, N83178, AB018288
1940	HTEEZ62	881052	۲.	, AI950251,
			present invention are one or more	, AI560075,
			polynucleotides comprising a	AA603342, AL135260, AW338106, AA505767,
			nucleotide sequence described by	AA888065, AI625041, AI909320, AI357213,
			the general formula of a-b, where a	AA962704, AI911938, H29506, AA353956, AI928495,
			is any integer between 1 to 1870 of	AA211037, AA581961, AI750915, AA516054,
			SEQ ID NO:1940, b is an integer of	AI750267, AA249644, AA211203, AI493165,
			15 to 1884, where both a and b	AW389552, AA104012, AI905441, AI887429,
	_		correspond to the positions of	
			TO.	AA182761, AI739109, AA182641, Z42725, AA638984,
			NO:1940, and where b is greater	AW389580, T48739, D19877, AA486796, AI697765,

			than or equal to a + 14.	AI300924,	AI873826,	N41871, A	AB020657, AI	AF161553,
				AJ012449,	AL078644,	AR018872,	AL137640	
1941	HOAAH52	881074	Preferably excluded from the	AI638708,	AW370588,	AA604391,	AI638200,	
			present invention are one or more	AL046090,	AI052244,	AW055067,	AW055206,	
	_		polynucleotides comprising a	AA224549,	AW375847,	AI679109,	AL042378,	
			nucleotide sequence described by	AI621228,	AW055056,	AI633697,	AW131512,	
			the general formula of a-b, where a	AI858264,	AI652500,	AA418385,	AW007559,	
			is any integer between 1 to 2717 of	AI347910,	AA633193,	AI417517,		
			SEQ ID NO:1941, b is an integer of	AL039518,	AI379655,	AI735776,	AI580118,	
			15 to 2731, where both a and b	AI611056,	AI767569,	AI332364,	AW006925,	
			correspond to the positions of	AA431974,	AI566498,	AA458620,	AI333573,	R93775,
			nucleotide residues shown in SEQ ID	AA633310,	AI804397,	AW190968,	AI304495,	
			NO:1941, and where b is greater	AW025852,	AI077447,	AI278898,	AA854076,	
			than or equal to a + 14.	AA400042,	AI081935,	H48411, A	H48411, AI061256, AI346015	1346015,
				AI042287,	AI200205,	AI298915,	AI150973,	
				AI400748,	AA705014,	AI921341,	AI206630,	
				AA258351,	AI493294,	AA418302,	W80672, AI378534	1378534,
			-	AI367993,	W80671, A	W80671, AI093517, AI445930,	1445930, A	AI307183,
				AA467763,	AA418344,		AA401498, AI267890,	
				AI953454,	AI271612,		N72284, AA937447, AA469431,	A469431,
				AI361498,	AI208143,	AA725419,	AA296397,	
				AA507583,	AA150850,		AI207267, AA865832, H18576,	H18576,
				AI056172,	W60546, H	13134, AI7	W60546, H13134, AI754190, AW338131,	38131,
				~	AI569024,	R69127, A	R69127, AA911897, AI028185,	I028185,
				N73581, R	80599, N91	387, H6319	R80599, N91387, H63197, AA232897,	7,
	·-				AA150542,		AI358148, AA921728,	A921728,
					AA132871, AI288107,		AA400712, AA742907,	A742907,
					AI290519, AI952567,		R11774, R68082	082,
				H60801, H	H60800, R69	246, T679C	R69246, T67909, T64951,	
				AI868438,			AI537951, AW235108,	W235108,
				AA232896,	N70399, AA342399,		T69432, H82789,	789,
				AA360349,				R80600,
				AI580686,	AA857394,	AI678572	AI678572, H18469, W	W04986,
				w	AA610546, H57599,	H57599, F	R80203, R91273	273,
				H57600, T	68057, H82	690, N753E	85	6,
				AL039517,	T52512, A	AL043057, R	R93722, N76405	405,

				AIS37427, AA400660	0, H82428, Z40015, H18502,
				AL044808, F04916,	R98833, AI474154, AI478
				AI934138, T96021,	AA133024,
				T54446, AA371002,	AL045017, R68119, T16415,
				AW271181, AA403235, AA676809,	5, AA676809, T70487, AA626926,
				R37695, F02870, H	R37695, F02870, H51082, R97530, AW389296,
				AA247471, AI93229	AA247471, AI932299, AW376391, Z44495, AW371130,
				R82536, AI933296,	AI933296, AL044806, AL043245, AI672519,
				AI133627, D87438	
1942	HSDXB50	881104	Preferably excluded from the	AI816760, AI346903	3, AI189171, AI860301,
			present invention are one or more	AA284405, AI340328,	8, AA485290, AW028742,
			polynucleotides comprising a	AW073309, AI539128,	8, AI749857, AA910895, N77735,
			nucleotide sequence described by		AI422690, AA868655, AA046578,
			mula of a-b,	AI038920, T32229,	T32229, AI936194, AA742438, AW001568,
			is any integer between 1 to 735 of		AW170086, W25066, AA296692, AI077505,
_			SEQ ID NO:1942, b is an integer of	AI375014, T95167,	
			15 to 749, where both a and b	AI348244, N36073,	
			correspond to the positions of	T63086, AI432379,	AA127847
			nucleotide residues shown in SEQ ID	AI082289, W31500,	
	•		NO:1942, and where b is greater	AI278762, T82102,	
			than or equal to a + 14.	AA877544, AA706829,	9, AI129303, AI361287,
				AW249798, AA594759,	9, AA524456, AA542925,
-				AI240209, AA126112,	2, AA934763, AI342601,
				AI052791, AI857321,	1, AI128632, AI340141,
				AW118892, N25202,	N25202, AA814658, AI041906, D11489,
				AA485295, AW00205	AW002059, AI370689, AA553675,
				AI609301, AI459183,	3, AA195893, AW088630,
				AIS61215, AI800091,	1, AW248136, AL050318,
				AF112213, S83364	
1943	HFKMJ24	881105	Preferably excluded from the	AA742438, AI346903	3, AW170086, AI816760,
			present invention are one or more	AI189171, AI43237	٥,
			polynucleotides comprising a	AW028742, AW07330	ο,
			equence	AI126547, AI749857,	7, N74204, AA910895, AI129303,
				AI038920, AI246120,	
			is any integer between 1 to 1208 of	AW249798, AA877544,	AI735203,

			SEO ID NO:1943, b is an integer of	AA868655,	AA542925,	AI375014,	AA934763,
			22, where both a	AI128632,	AI340141,	AW118892,	N92840, AI240209,
				AI348244,	AA706829,	N25202, AI346077,	I346077, AI342321,
			residue	AI748952,	AI857321,	AW002059,	AA553675,
			NO:1943, and where b is greater	AI052791,	AA127847,	AA814658,	AI041906,
			equal to a + 14.	AA983612,	AI609301,	AA994633,	AW006650,
				AI400295,	AA729483,	AI459183,	AA903651,
				AI800091,	AI561215,	H09610, AW088630,	W088630, AI683272,
				AI753574,	AI719306,	AI359224,	AI278762, T32229,
				AI819003,	AI093341,	D11489, A	AI342601, AW300745,
				AI374975,	AI346938,	AI183409,	AI423782,
	_			AA126006,	AA612604,	AA161217,	AA846503,
				AI284860,	AI275160,	N80744, H	H06158, AA844576,
				W16677, AI	I310420, AI539128,		AA996156, AA046578,
				AA737921,	AI985064,		N58366, AI827968,
				AA719050,	N26915, AI091923,		AI262701, AA524456,
				AI674584,	AA873274,	AI698929,	, AA485290,
				AA292533,	R99586, A	AI079471, A	AA806662, AI361287,
				T81787, A	AI370853, W.	W31500, AW1	AW193899, AI082289,
				AI805446,	AA583430, T58149,		H17502, F30305,
				AA594759,	W25066, AW248136,	W248136, A	AA195893, N77735,
				T95072, F3	30309, AA4	4	
				AW059835,	AW103745,	T95167, R	R35655, T82102,
				AI370689,	AA485295,	T23459, A	AW366963, AA564661,
				T63086, W	W40151, AA484058,		Κ
				AA126112,	AA296692,	_	
				AA192315,	AA911901,	N79525, A	AI784438, AW073849,
				AA913441,	AA534551,	T24804, A	AI074360, AW193751,
				H90230, A	F112213, A	AL050318, S	83364, AA689442
1944	неоосп	881219	Preferably excluded from the	AI924972,	AL046288,	AW189048,	W89124, AI091620,
			present invention are one or more	AA492579,	AA588728,	AI439428,	AA449355,
			polynucleotides comprising a	AA634228,	AI146362,	AA043859,	AA581516,
			nucleotide sequence described by	AA507328,	9	AA146720,	AI056656,
			formula of a-b, where	AA765659,	N64539, A	AL046287, A	AW402025, AA312475,
		_	is any integer between 1 to 2772 of	AI457992,	AW005493,	AA292416,	AA449614,
			SEQ ID NO:1944, b is an integer of	AA742592,	AA465004,	AA405756,	AA078819,

			15 to 2786, where both a and b	AA613822, N64732, AA405775, AA196964, AA367635,
			correspond to the positions of	AA373433, W88918, AA504065, AA652295, N91745,
			nucleotide residues shown in SEQ ID	T79620, AA996002, F25128; AI364464, AA515314,
			NO:1944, and where b is greater	AA394253, AA078918, AI909748, AA455284, N80334,
			than or equal to a + 14.	AL044772, AA377702, AA742682, AI583136,
				AI907986, AI909746, AA146721, T79705, AI798856,
				AA037697, H55648, AA
				AA814521, AI675619, AI872260, AW370721, R32993,
				D78805, D78848, AW078800, AW082532, AW020164,
				AI245304, AI688854, AI492648, AL096741,
				AC004882, AC005529, Z82171
1945	HWMB122	881221	Preferably excluded from the	AI800907, AI949684, AI052333, AW131568,
			present invention are one or more	AA732570, AA769120, AI743959, AI436302,
			polynucleotides comprising a	AW082175, AW273742, AI677956, AA037263,
			nucleotide sequence described by	AA885367, AA761521, AI936106, AI433128,
			the general formula of a-b, where a	AI292313, AI458263, AI687626, AI378687,
				AI187910, AI289598, AI378924, AI224510,
			SEQ ID NO:1945, b is an integer of	AI808484, AA890001, AI363454, AW340276,
			15 to 1483, where both a and b	AI077398, AI168640, W89211, W88447, AI566016,
			correspond to the positions of	AL043030, AA836573, AA768422, AA634503,
			യ	AI141297, AI539216, AA918633, AI350946,
				AA825685, AA515491, AA994089, AA609078,
			than or equal to a + 14.	AA761310, AI628981, AI206686, AW105192,
		_		AA776321, AA676705, AI676082, AA363995, D62240,
				AI094091, AI300249, AI400742, T98450, AI809452,
				N75907, U66469, U66471
1946	HETDL42	882330	Preferably excluded from the	AI344189, AI693945, N91690, AI457192, AW150901,
			present invention are one or more	AI798181, AA503831, AI458569, W86357, W86242,
			polynucleotides comprising a	N92074, T79381, W86600, AI915320, W90710,
			nucleotide sequence described by	R94236, AI282976, R94333, AA470366, T55160,
			the general formula of a-b, where a	H47818, T79811, W01906, N71011, A1702229,
			is any integer between 1 to 1573 of	
			SEQ ID NO:1946, b is an integer of	T61655, AA120932, AA579769, H24026, AW170681,
			15 to 1587, where both a and b	AI243696
			correspond to the positions of	AA345280, AI908519, AI051595, AA885499, W80464,
			nucleotide residues shown in SEQ ID	AA917596, AI380135, N29558, AI867394, AA250763,

NO:1946, and where b is greater	AI284328, AI803101, AW440273, AA603344,
 than or equal to a + 14.	AW148392, AA453747, H80554, AA453828, AA528253,
	W80573, AI254217, AW183037, AI419419, AI423034,
	AI305512, H65206, AA989137, AI559284, AI659077,
	LC)
	AW440223, AI073889, TS7089, AL046966, AI144070,
	AA962018, AA112330, AA630098, AI419982,
	AA954260, W93927, AW173728, R28013, AA146651,
	AI583416, AA668673, AA191610, F34079, AA703680,
	AA568394, AI053711, AW270496, AA069314,
	AI357477, AL041838, W02028, AA706521, AA664331,
	H89224, AW085628, AI207861, AI253208, AI744801,
	, AI769492, AI251385, AW2710
	493025,
	AA931216, AI991553, AI053773, AI311753,
	5022, AA
	AI252858, AI053963, AW086339, AA888155,
	AL135273, AI792443, AA083383, W92523, AI400721,
	AA504865, AW262442, AA789229, AI250275,
	AA011377, AI251700, AI254684, AI244896,
	, AW052205,
	AF050157, AL109654, AC005919, AC004062, U52112,
	AF030001, AC006289, AL132774, AL049636,
	, AC003949,
	, AL078630,
	, AP000338, AL031056,
	_
	AC004638, AF130342, AF084363, AF107258,
	AC003976, AC004551, AC002072, AC005619,
	AP000080, X79283, AF126403, AC003061, AC005972,
	AF095725, AC005921, AF052041, AL049780,
	AC004051, AC016026, AC005304, AF109905,
	AC007707, AF111103, AC005580, AL031864,
	AC006039, AC005740, AL022401, AC003107,
	AC006012, AC003664, AC006371, AC005587,
	AL031737, AF001549, AP000014, U85195, AC002470,

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	9	_			285987,
	œ	٦,	Z97987, AF	AF091512, X07	X07200,
	AC006387, A	AC004126, 1	AJ006996,	AC006525,	
	AL033533, A	AC007528, 1		AC006328,	
	AJ003147, A	AP000208, 1		X15051, AC0	AC005599,
	~		AP000247,	3,	U62317,
	X15052, ALO	AL022333, AC			AF139987,
	,				AP000952,
_	AF229844, Z	Z82203, AP	AP000039, AC	AC016025, U6605	,059,
		AF125314, 1	AC000116,	AC003694,	
	_	~	AC011331,	AC006370, Z	286062,
			AC004033,	AC005878,	
		7,	AC007277,	AL031010,	
	AL024509, A	AC006285, 1	AC005701,	AC008080,	
	AF131205, A	_	v	007425, AL1	AL121657,
	AC002080, A	_	AC009069,	AL031655,	
	AC000105, A	AC005881, A	AF130248, AC006368	AC006368,	
	_	Z82244, AL	AL031228, AC009396,		AC007115,
		_	AC007899,	AP000961,	
			AC006445,	AC002331,	
		_	AF064858,	AP000081,	
·-	_	_	AC006945,	AC005184,	
			AC007314,	AC005303,	
			AC007359,	AC004859,	
	_			AL035458,	
		_		AF196972,	
	_	AL049838, A	ς,	,	298748,
	_	AC004910, 2	Z82201, AC	AC008175, AL0	AL034412,
	AC005960, A	AC005553, 7	AC004848,	AL049631,	
	AP000697, AC	AC004217, #	AC008984,	AC006042,	
	AC006989, A	AF212831, 2	Z97054, AF027865,		AC006382,
	AC008033, AC	١,	AC007344,	AF060568,	
	3,	Z97353, AF1	_		AF107257,
	ω,	5216,	_	_	U91323,
	AB010266, Al	AL023582, A	AL034549,	AC007917,	

				AL049779, AP000313, AC009802, AC004467,
			•	
				AP001116, AL136363, AC004967, AL035684, AF034569
1947	HMEKW4	882715	Preferably excluded from the	AA553612, AA813301, 236965, D61366, AI216671,
	4		present invention are one or more	Z21245, AW152524, AI339525, AA483108, AI114701,
			polynucleotides comprising a	AI720301, AI375684, AI066646, AI755202,
			nucleotide sequence described by	AA584876, AA057530, AI341571, AW130427,
			the general formula of a-b, where a	AA584862, AW068996, AA569586, AW069783,
			is any integer between 1 to 1993 of	AA679937, Z86040, AC007385, AL031230, AC009247,
			SEQ ID NO:1947, b is an integer of	AB020874, AL049546, AL079304, AL021397,
			15 to 2007, where both a and b	AL035078, AC004890, AC004990, AC007103,
			correspond to the positions of	AC003009, AC004804, AL024498, AC004263,
			nucleotide residues shown in SEQ ID	AC005844, AL034375, AC005723, U91326, AC005409,
			NO:1947, and where b is greater	AL049539, AC006241, AC009509, AC007842,
			than or equal to a + 14.	AC006430, AL031296, AC005086, AC010205,
			,	AL023578, AC007528, AC006377, AC005081,
				AL021395, AC
				L133243,
				AL133245, AC004087, AL031684, AP000141,
				AC004821, AP000500, AC006478, Z93017, AC008372,
				AC004859, AC004125, AC006229, AC006525, Z78022,
				AL022576, AC004796, AL035249, AC005181,
				AC004028, AP001137, Z85986, AF045448, D87675,
				AL049696, AF001549, AC005670, U91318, AC005483,
				AR036572, U91328, AL049713, AC005180
1948	HCEDM42	882729	Preferably excluded from the	AI563939, AW250591, AA280100, AA148046,
			present invention are one or more	AI167949, AI160019, AA886389, AI679948,
			polynucleotides comprising a	AI523219, AA147993, W94919, AI679440, AA307127,
			nucleotide sequence described by	AA480164, N26434, R54543, AA064644, H08047,
			the general formula of a-b, where a	AI520745, H99329, R60593, R60646, AA064686,
			is any integer between 1 to 1236 of	AA283759, AA280033, R54445, AA303581, H07940,
			SEQ ID NO:1948, b is an integer of	AA283994,
			15 to 1250, where both a and b	AI085856, N70908, R11229, AI540673, AA809976,
			correspond to the positions of	AA909579, AA775556
			nucleotide residues shown in SEQ ID	
_			NO:1948, and where b is greater	

			than or equal to a + 14.	
1949	HCRNZ31	882762	Preferably excluded from the	AW388071, AW388070, AW392828, AW170095.
			present invention are one or more	AI139114, AA130783, AI796575, AI582280,
		_	o	5,
			nucleotide sequence described by	AI160038, AI631539, AI205291, AA143796,
			the general formula of a-b, where a	AA086002,
			teger between 1 to 2140	AW388098, AA086109, AI374885, AW392810,
			SEQ ID NO:1949, b is an integer of	AI146898,
			15 to 2154, where both a and b	AA303484, AI335908, AI917197, AI094414, W32500,
			correspond to the positions of	F02983, H77763, AA371674, D58760, AW131074,
			യ	AA148180, AW392820, AA148700, AA130888, R72708,
			NO:1949, and where b is greater	AW363332,
			than or equal to a + 14.	
				D59193, AA09
				AIS71527, AA151983, AA583490, F049
				AA303931, AA098988,
				AA148701, AA747401,
				AI479148, N28704, AW021399, W01939, AW270652,
				, AL042054, N71729,
				C004231, AC005971,
				AC005514, AC005527, AL022316, AC003980,
				AC007014, AL133245, AL117344, AC003950,
				AC004233, AP000229
1950	HWMBU8	883172	Preferably excluded from the	AA368362, T52098, R69052, R27072, AA397783,
	6		present invention are one or more	AA393589, T95399, AA912955, AW137196, AA155762,
			polynucleotides comprising a	AA188555
			nucleotide sequence described by	
			the general formula of a-b, where a	
			is any integer between 1 to 638 of	
			SEQ ID NO:1950, b is an integer of	
			15 to 652, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1950, and where b is greater	
			than or equal to a + 14.	
1951	HUFBY15	883201	Preferably excluded from the	AA625286, AA303053, AA303052, AA297581

re a of of of ID	M00425, AA349641, N42533, AI557558, AI557559, nore AW360991, R12333, AI557560, Z46216, AI890540, by lere a 11 of er of E	AL478843, AA628092, AI816845, AI813678, more AW269372, AI310217, AI742137, AI887196, AA722779, AA740417, AI363399, H94805, H95343, by AA890712, AA643210, AI743293, AI362725, nere a AI391652, AA410876, AI474205, AI261631, er of AI280434, AI832281, AW001746, AA449475, ar of AI459617, AW152661, W32215, H61131, AI190504, b AI282582, AI872611, W32179, AA449638, AI345648 E AI271086, AI473071, AJ245719 er	AA523290, AA700004, AI927220, AW170580, W74492
present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where is any integer between 1 to 455 of SEQ ID NO:1951, b is an integer of 15 to 469, where both a and b correspond to the positions of nucleotide residues shown in SEQ NO:1951, and where b is greater than or equal to a + 14.	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 741 of SEQ ID NO:1952, b is an integer of 15 to 755, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1952, and where b is greater than or equal to a + 14.	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 1008 of SEQ ID NO:1953, b is an integer of 15 to 1022, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1953, and where b is greater than or equal to a + 14.	Preferably excluded from the
·	883254	883371	883753
	нівсе91	HWLKF77	HOGCA75
	1952	1953	1954

	present invention are one or more	AI859845,	AI991311,	AA522795,	AI081052,
	polynucleotides comprising a	AA535079,	AI400364,	AI335984,	N
	eotid	AW170345,	AA622540,	AI273767,	AW168283,
	the general formula of a-b, where a	AI188508,	AA565989,	AI559433,	AI420481,
		AI246782,	AI928146,	AA157892,	AA314960,
		AI281336,	AW194453,	AA838633,	AA844471,
-	6, where	AI401064,	AI949231,	AI911649,	AI268908,
-	correspond to the positions of	AI874198,	AI186144,	AI819846,	AI276313,
	െ	AI874344,	AI963847,	AW193220,	AI863584,
	_	AW167101,	AW168206,	AA149417,	W79089, AA506616,
	than or equal to a + 14.	AIS64546,	AL036495,	AA434123,	AI560666,
		AA149738,	W02467, AA948146,		C06165, AI660464,
		AW167111,	AI961910,		AW194388,
		AI567796,	AW009339,	AW009339, AA434059, AI739607	AI739607,
		AI280032,	R48300, AA551656,	4551656, AI	AW167849, AI346572,
		AI923100,	AI005290,	AI091394,	
		AI588982,	AI819915,	AI950029,	ın
		AI347074,	AI347076,	AI660868,	AW374558,
		AI682624,	AI348165,	AI949885,	AI347071,
		AW014104,	AA582757,	AI860565,	AI222884,
-		AI861959,	AI283186,	AI347501,	AI305833,
		AI031766,	AI346386,	AI346944,	AW189088,
		AI032425,	AI283162,	AI347072,	H27323, AI214245,
		AI346606,	AI743195,	AW015201,	AI347060,
		AI346569,	AW275383,	AI281140,	AI346475,
		AI743978,	AI274133,	AI738882,	AI273374,
-		AI347930,	AI738627,	AI991114,	AI097004,
		AI144005,	AI304544,	AA569935,	AI281141, U46417,
		AA157596,	AI274318,	AI285074,	AI346274,
		AI336454,	AI346908,	AW374542,	AI339875,
		AI014860,	AA293207,	AI339827,	AI861957,
		AI281257,	AI243957,	AI281300,	AI336446,
		AI660830,	AI347929,	AI368165,	AA477634,
		AA411444,	AI343934,	AI636236,	AI274312,
		AI424819,	AW024873,	AI337303,	AI339815,
		AI470046,	AI690641,	AI284953,	AI284985,

1955	HOGCJ47	883799	Preferably excluded from the	AW054994, F33829,		A1560717, A	AI268302, A	AW005178.
			present invention are one or more	F22745, AA284546,				AI356840,
			polynucleotides comprising a	AI493477, F36987, AI081004,	F36987, A			AI633219,
			nucleotide sequence described by	T66954, T3	6169, W71	T36169, W71988, Z39991, H50924	1, H50924,	
			the general formula of a-b, where a	_	F09164, AA043299,	A043299, T	T31835, M78780,	780,
		_	is any integer between 1 to 1115 of		H16657, A	H16657, AW262658, AA745578, AA744099,	A745578, A	A744099,
			SEQ ID NO:1955, b is an integer of	AI349099,	AA989269,	AA989269, R72575, H51586, AA74439	51586, AA7	44396,
			15 to 1129, where both a and b	T79883, W7	6380, H16	T79883, W76380, H16514, H38527, AA995198,	7, AA99519	8,
			correspond to the positions of	AA296888,	AA541441,	F11503, A	I475083, A	AI302606,
			nucleotide residues shown in SEQ ID	AA043300,	AA886838,	R54219, A	AI125823, T	T66953,
					AW361009,	AW361009, AA296951,	F03443,	A297044,
		•	than or equal to a + 14.	AA335686,	F05047, R	F05047, R37601, AA090754, AI970619	90754, AI9'	70619,
				Z44304, AW	374215, A	Z44304, AW374215, AI547101, R51823, AA783044,	51823, AA7	83044,
	_			AA594940,	AW176749,	AW176749, AA583598,	T15585, R	R49122,
				AA085248,	AF131774			
1956	HWLUT61	883945	Preferably excluded from the	AI942421,	AA588562,	AI942402,	AI520886,	
			present invention are one or more	AI867203,	AA995170,		AW380270,	
			polynucleotides comprising a	AI680440,	AI362487,		R82350, A	AI934005,
			nucleotide sequence described by	AW089784,	C04722, A	1046708,		AA016994,
			the general formula of a-b, where a	AI274637,	AI872632,	D19775, A		AL049685,
			is any integer between 1 to 265 of	AL049792,	AF093744			
	-		SEQ ID NO:1956, b is an integer of					
			15 to 279, where both a and b					
			correspond to the positions of					
			·O					
			NO:1956, and where b is greater					
			than or equal to a + 14.					
1957	HLTBA42	883971	<i>'</i> α'	AI767559, 1	AI631820,	AI758931,	AI758389,	
			present invention are one or more		AA630485,	AA761469,	AW195693,	T89742,
			polynucleotides comprising a	AA807177, 1	AA361233,	AI679708,	AI244041,	
			nucleotide sequence described by	AI572549, 1	AA947977,	AI679134		
			the general formula of a-b, where a					
			is any integer between 1 to 909 of					
			SEQ ID NO:1957, b is an integer of					
			15 to 923, where both a and b					
			correspond to the positions of					

			nucleotide residues shown in SEQ ID NO:1957, and where b is greater than or equal to a + 14.	
8561	нненв82	884038	Preferably excluded from the present invention are one or more polynucleotides comprising a	AI676130, AI991800, AI936232, AA307685, W67860, AI640485, AI628790, AA524353, AI824956, AI890762, AI800990, AI335005, N31143, N21294,
			nucleotide sequence described by the general formula of a-b, where a	, AW302169, AW002644, , AA983846, AA595031,
				H-
			15 to 1757, where both a and b	AAS79237, AW015641, T64746, T31944, AAS70191,
			correspond to the positions of nucleotide residues shown in SEO ID	AA084445, AA935035, CIS927, AA358195, AA081627, T07826, N94623, T34036, Z44938, AA380035,
				-
			than or equal to a + 14.	AW089
				AW373174, AW373195, W23822, Z41687, T83793,
				AA358196, AI360228, AA129234, AA913439,
				AF090992
1959	HE2PR08	884095	Preferably excluded from the	AA582960, AI
			present invention are one or more	AA224518, AA524291,
			polynucleotides comprising a	AW089837, AA846846,
			nucleotide sequence described by	, AI798847, AA582557,
			the general formula of a-b, where a	, AI809936, AW173427,
			is any integer between 1 to 2842 of	_
			SEQ ID NO:1959, b is an integer of	AI2438I8, K80863, AA/3638/, AII6/988, I//889,
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	AA323841,
			NO:1959, and where b is greater	T77712, A1912397, AW117749, AW173596, AA653386,
			than or equal to a + 14.	85
				7
				4634686, AA091136
1960	HMKAN71	884161	Preferably excluded from the	AI635715, AW411210, AI624534, AA879465,
			present invention are one or more	AW104990, AW409582, AI766309, AA081177,

			polynucleotides comprising a	AI803484,	R78080, AI	AI129966, AI925109	.925109, AI804159,
			nucleotide sequence described by	AA279212,	AA410910,	AA678827,	AI860837,
			the general formula of a-b, where a	AI183591,	AW316983, AI431314,	AI431314,	AA766602,
			is any integer between 1 to 1706 of	AA081236,	AW194027, AI521521,	AI521521,	Z38832, AA588351,
			SEQ ID NO:1960, b is an integer of	AI923638,	N39554, R22273, AA447188,	2273, AA44	7188, AA769352,
			15 to 1720, where both a and b	T52102, A	T52102, AA371263, AA259257, T60532,	259257, T6	0532, AW411209,
			correspond to the positions of	R22218, Z	Z42670, AA443811, AA969814,	3811, AA96	39814, AA729654,
			nucleotide residues shown in SEQ ID	AA259256,	59256, AI969030, AW409826,	AW409826,	R24524
			NO:1960, and where b is greater				
			than or equal to a + 14.				
1961	HSIFV30	884168	Preferably excluded from the	AI660957,	AW361534,	AW361532,	AI802756,
			present invention are one or more	AW361521,	AW361520,	AW009763,	AI660234,
			polynucleotides comprising a	AI802693,	AW361523,	AI721275,	AA581198,
			nucleotide sequence described by	AW361522,	AW361528,	AA296955,	AI721121,
			the general formula of a-b, where a	AA508854,	AA297150,	AW009764,	D25727, AI687981,
			is any integer between 1 to 2840 of	AI582072,	AF127036,	AF039400,	AF095584,
			SEQ ID NO:1961, b is an integer of	AB017156,	AF039401,	195746	
			15 to 2854, where both a and b				
			correspond to the positions of				
			nucleotide residues shown in SEQ ID				
			NO:1961, and where b is greater				
			than or equal to a + 14.				
1962	HNTSY52	884215	Preferably excluded from the	AI815240,	AI631739,	AA309645,	AI696961,
			present invention are one or more	AI479235,	AA307961,	AI978872,	AW195761,
			polynucleotides comprising a	AA280818,	AI990440,	AW262762,	AI809185,
			nucleotide sequence described by	AI037930,	AI637988, 1	AI754009,	AA181165,
			the general formula of a-b, where a	AA972531,	AI817057, 1	AI494056,	AW073248,
			is any integer between 1 to 4073 of	AA181166,	AI826853, 1	AI361369,	AI149286,
			SEQ ID NO:1962, b is an integer of	AI752584,	W52618, AW339206, AW075435,	339206, AW	1075435, AA115631,
			15 to 4087, where both a and b	AI445241,	AI523220, M62298, AA558913,	M62298, AA	558913, AW368570,
			correspond to the positions of	NS1760, A	N51760, AA348679, AI735744, AW384980,	735744, AW	1384980, AW384967,
			nucleotide residues shown in SEQ ID	AI802541,	Z19223, N3	5007, N741	Z19223, N35007, N74118, H03102,
			NO:1962, and where b is greater	AA102848,	Z25028, AI	624448, AI	Z25028, AI624448, AI279412, AI476071,
			than or equal to a + 14.	AA385867,	AA095022, AW194583, AI383593,	AW194583,	AI383593,
				AA360919,	R79669, Z28444, AA506352,	8444, AAS0	6352, R26853,
				AA133388,	AA330074, 1	, N30413, Z2	Z28730, AA020013,

				AI954282, R79858, D31597, R77935, AA280996,
				H99307, AA020014, R27081, AI950631, AA295264, AA402581, AA093272, AA093324, AA248050,
1963	HCROM43	884379	Preferably excluded from the	
			present invention are one or more	T09212, T31698, T83796, AA714176, T27030,
			polynucleotides comprising a	AI655004, AW239098, AF196972
			nucleotide sequence described by	
			is any integer between 1 to 787 of	
			SEQ ID NO:1963, b is an integer of	
			15 to 801, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1963, and where b is greater	-
			w	
1964	HLWCF60	884529		AI083497, H14688, N77514, AW015613, H16869,
			present invention are one or more	AA377154, AW194949, AA378912, AW390260, H24407,
			polynucleotides comprising a	AA307120, W39491, F25064, AA252725, AI539349,
			nucleotide sequence described by	
-			the general formula of a-b, where a	R57305, H06942, AA488566, AF151908
			is any integer between 1 to 1612 of	
			SEQ ID NO:1964, b is an integer of	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1964, and where b is greater	
			than or equal to a + 14.	-
1965	HWLKD85	884719	1	AA121115, AA323118,
			present invention are one or more	AA325395, AA248006, AB028859, AJ250137
			polynucleotides comprising a	
			nucleotide sequence described by	
			the general formula of a-b, where a	
			is any integer between 1 to 576 of	
			inte	
			15 to 590, where both a and b	

			correspond to the positions of					Γ
			nucleotide residues shown in SEQ ID					
			NO:1965, and where b is greater					_
			than or equal to a + 14.					
9961	HCRMX54	885350	Preferably excluded from the	AL038837,	AL037051,	AL039074,	AL039128,	<u> </u>
			present invention are one or more	AL039109,	AL039108,	AL039659,	AL039156,	
			polynucleotides comprising a	AL045337,	AL039625,	AL039648,	AL039629,	
			nucleotide sequence described by	AL039678,	AL042909,	AL040992,	AL039564,	
			the general formula of a-b, where a	AL038531,	AL037726,	AL045353,	AL036973,	
			is any integer between 1 to 1956 of	AL044407,	AL039410,	AL039423,	AL039538,	_
			SEQ ID NO:1966, b is an integer of	AL039386,	AL044530,	AL039566,	AL039509,	•
			15 to 1970, where both a and b	AL036725,	AL045341,	AL039150,	AL036196,	
			correspond to the positions of	AL037639,	AL038025,	AL039924,	AL036767,	
			O.	AL037615,	AL038821,	AL036117,	AL036238,	
			NO:1966, and where b is greater	AL043441,	AL045794,	AL039085,	T24119, T24112,	
			than or equal to a + 14.	AL036679,	AW013814,	AL043445,	AL043422,	
		·		AL037526,	AL037027,	AL037601,	AL043423,	
				AL036924,	AL036964,	AL036158,	AL036765, H00069,	
				AL036268,	AL036733,	AL037177,		
				AL036418,	T23947, AI	1036998, TO	T23947, AL036998, T02921, AL036133,	-
				AW451070,	AL037643,	AL036132,	AL037082,	
				AL038851,	AL036167,	AL036163,	AL037178,	
				AL037049,	AL037085,	AL036190,	AL037600,	
				AL036914,	AL036139,	AL037047,	AI535983,	_
				AL037124,	AI535783,	AI535783, AL037021,	AL036191,	
				AW452756,	Z99396, AI	Z99396, AL044960, AL036152,	.036152, R47228,	
				AL036900,	D51250, AI	D51250, AL036150, AL036227,	.036227, AL048425,	
				AL036207,	AL036174,	AL036174, AL036953, AL036719	AL036719,	
				AL037679,	T23659, D8	D80253, AL036858,	6858, AL037077,	
		-		AL036808,	D59787, AI	.038043, AL	D59787, AL038043, AL037569, D80043,	
				DS9275, D	30219, T485	D59275, D80219, T48598, AA514190,	.90, 225782,	
				AL038447,		D80227, AW450376, D80240,	10240, D80134,	
				AA631969,	AL037002, D51423,	D51423, T1	T11051, AL036999,	
				D80210, Z	35783, D596	519, H00072	Z25783, D59619, H00072, AL037016,	
				C14227, AI	037094,	.036630, D8	D80193, D80196,	
				AW135155,	D80168,	AL039440, D5	D59927, AIS57751,	

	AL036229, AL039076, AL037742, D80366, AL043868,
	AW392670, AL038509, AL039077, AL119457,
	AL119324, AI142134, AL042544, C75259, AW451416,
	AL038520,
	AW372827, AW363220, AL119497, AL119319,
	AL119355, AL119483, AL119363, U46349, AL119391,
	AL119484, C14389, U46341, U46350, AL119522,
	AL119418, U46351, AL119341, AL119335, AL039504,
•	AL039555, AL039521, AL119396, AL039476,
	AL043586, AL044412, AL044364, U46346, AL119496,
	AL037205, AL119439, AL042984, AL119464,
	AL134527, AL134538, AL042614, AL042965,
	AL042975, AL043029, U46345, Z96142, V00745,
	X73004, AR036903, E13740, I19517, A76773,
	A22413, I13349, A11245, A35536, A35537, A02135,
	A02136, A10361, A04663, A04664, I08051,
	AF118808, I01992, A92636, E03165, E02221,
	E01614, E13364, X68127, A95051, AR062871,
	AR031374, A49700, AR031375, A58521, AR020969,
	', AR017907, AR036905, A382
	I56772, I95540, AR018924, A63067, A51047,
	A63064, AR018923, A48774, A63072, A48775,
	AR015961, A85477, AR035975, AJ244003, AJ244004,
	177, A85396, AR03
	A98767, I19516, A93963,
	A02712, I60241, I60242, A95052, AR043602,
	AR043603, AR043601, A95117, A18053, I06859,
	A84772, A84776, A8
	A84775, AR062872, A84774, AR062873, AR067731,
	AR037157, AR067732, A86792, A58522, A91750,
	A24782, A81878, I03343, AR022240, A97211,

A13392, A13393, A27396, I21869, A49045, E16678, D28584, I25027, I26929, I26927, I26927, I26929, I26927, I26926, A91753, I00079, E16590, A023093, A91965, A67226, A20701, A04710, Y11926, I03665, I03664, D88984, I66498, I66497, I66496, AR038286, I25041, I92488, AF156303, AR028564, AR06A957, I4698, A16598, A18755, A25856, A97221, AF019720, AF15638, A14595, A18755, A25856, A97221, AF019720, AF15638, A60990, A60987, D44443, Y17188, A10363, A1449, AF096793, Y17188, A10363, A1449, AR03312, I07888, A1672548, A1637672, AA02312, I07886, W05306, AA029735, AA3316, Integer of and booms of nin SEQ ID greater				A02710, E12615, AR035193, E14304, A07700.
121869, A49045, E16679, B26297, B26292, B262				A13393, A27396, AR027100, I2826
126927, 126929, 126929, 126927, 126929, 126927, 126929, 126927, 126927, 126927, 126927, 126927, 12692, 12		•		869, A49045, E16678, E16636, A82653,
126527, A58525, A70040, A1753 100079, A17515 100079, A1753 100079, A1753 100079, A1753 100079, A1753 100079, A1753 100079, A17526, I013664, D88944, I03664, D88944, I03664, D88944, A1710, X11926, A17526, A17				584, I25027, I26929, I44515,
HTPHK88 885476 Preferably excluded from the polynucleotide sequence described by the general formula of a by where a is any integer between 1 to 1208 of sequence control of the positions of nucleotide residues shown in SEQ ID NO.1967, bis an integer of correspond to the positions of nucleotide residues shown in SEQ ID NO.1967, and where be prefered to the positions of nucleotide residues shown in SEQ ID NO.1967, and where be prefered to the positions of nucleotide residues shown in SEQ ID NO.1967, and where be prefered to the positions of nucleotide residues shown in SEQ ID NO.1967, and where be prefered to the positions of nucleotide residues shown in SEQ ID NO.1967, and where be prefered to the positions of the positions are propertions.				7, A58525,
A1755, 100709, E16590, A20701, A07101, A0710				I44516,
### AJ19093, #91965, #6722 ### AJ19093, #91965, #6722 ### AJ19093, #91965, #6722 ### AJ19093, #91965, #91926, 10366, 10366, 108989, 106496, #91964 ### AJ19093, #91966, 10366				A91753, I00079, E16590, AF156294, AJ244005,
### ### ##############################				3, A91965, A67220, Y11923, AR02706
103665, 103664, D88984,				A04710,
HTPHK88 885476 Preferably excluded from the persent invention are one or more polynucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 1208 of sequence described by correspond to the postinons of morleotide residues shown in SEQ ID NO:1967, and where b is greater.				I03664, D88984, U87250,
HTPHK88 885476 Preferably excluded from the polynucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 1208 of 15 to 1222, where both a and b correspond to the positions of a uncleotide residues shown in SEQ ID NO:1967, and where b is greater. AR08286, 125041, 19248; AR08564, AR08569, AR08569, AR0870, D14548, AR08512, AR08513, AR08516, AR08513, A				
HTPHK88 885476 Preferably excluded from the polynucleotides comprising a polynucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 1208 of sequence described by correspond to the positions of nucleotide residues shown in SEQ ID NO:1967, and where b is graved by sequence between 1 sequence described by correspond to the positions of nucleotide residues shown in SEQ ID NO:1967, and where b is graved by sequence described by correspond to the positions of nucleotide residues shown in SEQ ID NO:1967, and where b is greater				
HTPHK88 865476 Preferably excluded from the polynucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 1208 of SEQ ID 15073, A00393,	_			
HTPHK88 885476 Preferably excluded from the polynucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 1208 of SEQ ID NO:1967, and where b is greater				A0
HTPHK88 885476 Preferably excluded from the polynomic sequence described by the general formula of a-b, where a is any integer between 1 to 1208 of SEQ ID NO:1967, b is an integer of nucleotide residues shown in SEQ ID NO:1967, and where b is are		*		X13220,
HTPHK88 885476 Preferably excluded from the polynucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 1208 of SEQ ID NO:1967, and where bis genetar is greater.				A18755, A25856,
HTPHK88 885476 Preferably excluded from the polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 1208 of SEQ ID NO:1967, and where b is qreater.				Ą
HTPHK88 885476 Preferably excluded from the prosper time general formula of a-b, where a is any integer between 1 to 1208 of SEQ ID NO:1967, b is an integer of nucleotide residues shown in SEQ ID NO:1967, and where b is qreater NO:1967, and where h is quarant NO:1967, and where h is qreater NO:1967, and where h is quarant NO:1967, and where h is qreater NO:1967, and where h is quarant NO:1967, and where h is qreater NO:1967, and where NO:1967, and where h is qreater NO:1967, and where NO:1967, and wher				AF156304, A91754, M32676, AB012117, AF096810,
HTPHK88 885476 Preferably excluded from the polynucleotide sequence described by the general formula of a nucleotide residues both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1967, and where b is a greater				I69350, S65373, X58217,
HTPHK88 885476 Preferably excluded from the polynucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 1208 of SEQ ID NO:1967, b is an integer of nucleotide residues shown in SEQ ID NO:1967, and where b is greater				, A60957,
HTPHK88 885476 Preferably excluded from the polynucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 1208 of SEQ ID NO:1967, b is an integer of nucleotide residues shown in SEQ ID NO:1967, and where b is greater				Y11449, AF096793, AR0664
HTPHK88 885476 Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 1208 of SEQ ID NO:1967, b is an integer of nucleotide residues shown in SEQ ID NO:1967, and where b is greater				A60987,
HTPHK88 885476 Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 1208 of SEQ ID NO:1967, b is an integer of nucleotide residues shown in SEQ ID NO:1967, and where b is greater				, A10363, AF130655, X7300;
HTPHK88 885476 Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 1208 of SEQ ID NO:1967, b is an integer of 15 to 1222, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1967, and where b is greater				, E04616, I03663, I(
HIPHK88 885476 Preferably excluded from the present invention are one or more present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 1208 of SEQ ID NO:1967, b is an integer of 15 to 1222, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1967, and where b is greater	+			AR063812, I07888, Y11920
ore W05306, AA029735, AA331672, W93893, A1672548, A1637672, AA025077, R26502 by ere a 08 of r of EQ ID		885476	0	AA433834, AA427986, W38581, AA362763, AA331674,
by ere a 08 of r of EQ ID			present invention are one or more	AA029735, AA331672,
by os r c			otides comprising	AI637672, AA025077,
ere os r r			e sednence	
7 C C L			of a-b, where	
r o r			en 1 to 1208	
r EQ			ы	
r EQ			15 to 1222, where both a and b	
shown in SEQ			correspond to the positions of	
, and where b is			shown in SEQ	
			, and where	

			than or equal to a + 14.	
1968	нсоврзя	885484	Preferably excluded from the	N55045, AI
			present invention are one or more	AC004388, AC004993, AC010722, AC006924,
			polynucleotides comprising a	AL033397, AL022151, Z84720, AL109654, AC005145,
			nucleotide sequence described by	AL136297, AC004081, AL121823, AC007458,
			the general formula of a-b, where a	AP000493, AC005053, Z93403, L11910, Z72001,
			is any integer between 1 to 1424 of	AL121654,
			SEQ ID NO:1968, b is an integer of	AL034377,
			15 to 1438, where both a and b	AL049588,
			correspond to the positions of	AL031650, AL117667, Z83848, AC003080, AC005250
			nucleotide residues shown in SEQ ID	
			NO:1968, and where b is greater	
			than or equal to a + 14.	
6961	HLQFI67	885511	Preferably excluded from the	Η.
			present invention are one or more	AW341505, AI590115, AI884695, AI651965,
			polynucleotides comprising a	AI863337, AI028587, AI246696, AI920847, R76087,
			nucleotide sequence described by	AI032590, AA835680, AA508647, AA765513,
		_	the general formula of a-b, where a	AI791278, H51121, AI568523, AA034147, AA513202,
			is any integer between 1 to 509 of	AA053714, T99214, AI821534, Z82198, Z82201,
			SEQ ID NO:1969, b is an integer of	AC008014, AC005296, AL031782, AL133512, Z74696,
			15 to 523, where both a and b	Ų
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	AC004836, AC005069, AC004068, AL049648, U69569,
			NO:1969, and where b is greater	AC006325, AC006256, AC007126, AC004106,
			than or equal to a + 14.	AF093117, AL049828, AL023806, AC002078, Z72004,
				AL049734, AC005066, AC006406, AL023582,
				AC006368, Z70288, AL133246, AC008080, AF165175,
				AC007370, AC005539, AC007461, AC005738,
				AL023579, AL022477, AL035684, AL022576,
				AC002526, AC007542, AL132800, AF165176,
				AL078598, AC008126, AC008072, AF064860,
				AL031681, AC007385, AC005232, AC004885,
				_
				AC003046, AL035686, AC007016, AL078602,
				AL109612, AL117355, U85197, AJ010598, AL135746,
				AC006143, AC006032, AL035667, AP000243,

				AP000203, AL034417, AF042090, U71148, AC005533, AC004042, AL079352, AL049844, AL031123,
1970	HAJBV26	886331	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 761 of SEQ ID NO:1970, b is an integer of 15 to 775, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1970, and where b is greater than or equal to a + 14.	AW392670, AL119483, U46341, AW372827, AW AL119484, AL119363, AL U46350, U46347, U463 AL119396, U46346, AL AL134533, AL119439, AL042970, AL134538, AL134518, AL037205, AL042995, AL134531, AL042995, AL134531, AL042995, AL13464, AL043003, AL043019, AL043003, AL119464, AR069079, U27699
	нвль90	886505	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 1120 of SEQ ID NO:1971, b is an integer of 15 to 1134, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1971, and where b is greater than or equal to a + 14.	AI291206, AI692352, AA159669, AA166774, W87878, H60270, R00390, AI174957, AA082398, AA047213, AI567717, N58610, AA384188, AA344124, AI970562, AI572002, AI860354, AA035047, N26366, AA382178, R21443, AA649513, AA294966, AA393451, AW372027, AW383791, N79097, AW176696, AA579377, AW383795, AW363037, AW372042, AW372015, AI887591, AW383956, AI590368, AA489105, AW363951, AA047214, AW372040, AA989009, AA286892, AW363951, AA047214, AW372040, AA459578, AW383793, AW3833800, AA092369, AW383794, AW364575, AW383786, AC004686, AF161410
1972	HWLFB44	886527	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 437 of	AI688604, AI660552, AI659950, AW296326, AW291582, AI700219, AI380340, AW004785, AW295479, AW006764, AI688540, AA522452, AA594441, AI695451, AA470898, AA594533, AIS81787, AI581803, AI581880, AI832419

			SEQ ID NO:1972, b is an integer of 15 to 451, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1972, and where b is greater than or equal to a + 14.	
1973	HCE4U96	886788	preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 1371 of SEQ ID NO:1973, b is an integer of 15 to 1385, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1973, and where b is greater than or equal to a + 14.	, T09220, AA338971, AI969431, AI8, Z42464, W46479, AW163719, AW1399, AI214207, AC004382
1974	HWLEL48	886914	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 734 of SEQ ID NO:1974, b is an integer of 15 to 748, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1974, and where b is greater than or equal to a + 14.	, AW376283, I82554, U7
1975	HTGBT14	887098	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 757 of	AA528172, AI870515, AW022634, AI122636, AI807139, AI524135, AW117562, AI332968, W94241, AI034051, AW119174, N53839, AI378914, AI708759, AA699609, AA425884, AA909771, AI086409, AI312652, AI382156, AI161356, AA635388, AA633491, W94238, W46444, AA746370, AA228039,

	SEQ ID NO:1975, b is an integer of	9, AA975136, AI144548, W9411
	15 to 771, where both a and b	.1350918, AI301665, AA92
	correspond to the positions of	AA702159, AI052284, AI340996, W
	res	AA228149, AI497988, AA084519, AA223979, F22291,
	NO:1975, and where b is greater	W262545, AI421254, W69785
	equal to a + 14.	
	•	
		AI718892, AA978346, H51405, AA866163, N73336,
		T48735, F26124, AI971845, W78797, AA704978,
		AA002051, AA463446, AA970170, W95702, F36672,
		F20308, R33196, AI460269, F34207, W95701,
		0
		F18648, AA428745, AA093730, AA666150, AA062817,
		AI027170, AA001847, AI264217, AI653972,
-		8
		AI269862, AI364788, AL047763, AL041150,
		AL042628, AW198075, AI537989, AI932794,
		, AI334450,
		AI648663, AI344928, AI358701, AI582932,
		AL036638, AL045500, AI570807, AL045266,
		AW079572, AI308032, AI698391, AI344785,
		5, AI889953,
		AA225339, AI345148, AI433976, AL037454,
-		AI620284, AI468872, AW020693, AI335209,
		, AI270183, AI554821, AW1511
		AI539771, AI537677, AI494201, AI802542,
		AI500659, AL036631, AW168485, F27788, AI815232,
		AI801325, AI500523, AI866090, N80094, AI923989,
		AI284517, AI500706, AI445237, AI491776,
		AW151138, AI889189, AI521560, AI500662,
		AI284509, AI288285, AI889168, AI866573,
	-	, AI627988,
		05769, AI888661, AI284513, AI88811
_		AI524671, AW162194, AI889147, AI812015,

AI440252, AI306613, AW051088, AI433037,
AI632408, AI886181, AW268302, AA715307,
 AW072719, AI933589, AI611348, AI635067,
AI610645, AL040243, AW103371, AI608936,
AI874166, AI254731, AI921248, AI819976,
AW023859, AL119791, AL043981, AI886753,
A1349004, A1686906, A1927755, AL121270,
AI798456, AW051258, AL042551, AI624293,
AI611738, AW148970, AI571909, AI619502,
 AI677796, AI352497, AI349598, AI684021,
AI288305, AW118518, AL039276, AW269097,
AW026882, AI923370, AI269205, AI064830,
AI929108, AI436429, AW193125, AL110402,
AI371228, AI500061, AI572892, AI613548,
AW083804, AI654276, AI620089, AC004985,
AF161453, AF015416, A12297, I89947, AL133014,
 AL137271, AL122049, AF111851, AF091084,
AF118094, AL133072, A08913, AF078844, AL137521,
6, AF026816,
F008439, A08910, I
AL050138, X72889,
AL137459, U80742, AF090901, X98834, AL049464,
AF106862, U72620, AL122110, AR011880, AL133080,
, AF125949, A65341, Z82022, AF09
3, AF087943, AL1336
103321, AL137560, AL117460, Y14314, AL050149,
4, E07361
AL110196, A77033
A58524, A58523, I00734, X93495, X65873,
AF113690, AF090934, AF113677, Y11254, AL049382,
3, AL050277, AL050116
7, E00778, AL122093, AL050393, AL122
A08912, I26207, AF104032, AF067728, U00763,

y excluded from the nvention are one or more or corprising a le sequence described by al formula of a-b, where a				AJ238278, X63574, AJ012755, AL122123, AL133104,
HKLRB09 887114 Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a				AF017437, AF097996, AL050024, AL133640,
HKLRB09 887114 Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a				
HKLRB09 887114 Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a				×
HKLRB09 887114 Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a				Y11587, AL137550, AF090943, AF183393, AF158248,
HKLRB09 887114 Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a				AL137292, S61953, U67958, I42402, A93350,
HKLRB09 887114 Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a				AF017152
HKLRB09 887114 Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a				4,
HKLRB09 887114 Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a				8, AL110221,
HKLRB09 887114 Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a				S,
HKLRB09 887114 Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a				5, AF113691, E07108,
HKLRB09 887114 Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a				6, AL137526,
HKLRB09 887114 Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a				8, AJ242859, L31396,
HKLRB09 887114 Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a				L31397, AI
HKLRB09 887114 Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a				5, AF057300,
HKLRB09 887114 Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a				
HKLRB09 887114 Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a				, A90832, AI
HKLRB09 887114 Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a				
HKLRB09 887114 Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a				
HKLRB09 887114 Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a				M30514, X84990, AL080127, AL
HKLRB09 887114 Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a				
HKLRB09 887114 Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a				
HKLRB09 887114 Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a				E04233, U96683, AL133568, AJ006417, X53587,
HKLRB09 887114 Preferably excluded from the present invention are one or more polynucleotides comprising a AL039109 nucleotide sequence described by the general formula of a-b, where a AL039648				AR013797, AL133081, AL1101
HKLRB09 887114 Preferably excluded from the present invention are one or more polynucleotides comprising a AL039109 nucleotide sequence described by the general formula of a-b, where a AL039648				3, U68387, AL137523, X87582, US
HKLRB09 887114 Preferably excluded from the present invention are one or more polynucleotides comprising a AL039109 nucleotide sequence described by the general formula of a-b, where a AL039648				AF081195,
HKLRB09 887114 Preferably excluded from the AF081197 present invention are one or more AW138645 polynucleotides comprising a AL039109 nucleotide sequence described by the general formula of a-b, where a AL039648				
HKLRB09 887114 Preferably excluded from the AF081197, U49908, present invention are one or more AW138645, AL038837 polynucleotides comprising a AL039109, AL039108 nucleotide sequence described by AL038531, AL039655 the general formula of a-b, where a AL039648, AL039625				E08631, Y10080, L19437, I09499
HKLRB09 887114 Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a AL039648, AL039629, AL039678 AL039648, AL039629, AL039678				X92070, AL137705,
HKLRB09 887114 Preferably excluded from the AI732659, AI791955, present invention are one or more AW138645, AL038837, polynucleotides comprising a nucleotide sequence described by AL039519, the general formula of a-b, where a AL039648, AL039629,	\dashv			U49908, AL
nvention are one or more AW138645, AL038837, otides comprising a AL039109, AL039108, e sequence described by AL038531, AL039659, al formula of a-b, where a AL039648, AL039629,		887114	$^{\circ}$	AI791955, AA577625,
otides comprising a AL039109, AL039108, e sequence described by AL038531, AL039659, al formula of a-b, where a AL039648, AL039629,			present invention are one or more	5, AL038837, AL039074,
e sequence described by AL038531, AL039659, al formula of a-b, where a AL039648, AL039629,			otides comp	AL039108,
al formula of a-b, where a AL039648, AL039629,			e sednence	AL039659,
			general formula of a-b, where	, AL039629,
teger betwe			teger between 1 to 1698	0, AL039128,

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	SEQ ID NO:1976, b is an integer of	AL042909,	AL039423, AL039410, AL039085,
	15 to 1712, where both a and b	AL045353,	AL036973, AL044407, AL039538,
	to the	AL039924,	AL039386, AL038821, AL044530,
	nucleotide residues shown in SEQ ID	AL039566,	AL039509, AL036196, AL043445,
	and where b is great	AL037526,	AL037639, AL038025, AL036418,
	equal to a + 14.	AL045341,	T24119, AL043422, T24112, AL037615,
		AL036767,	AW013814, AL043441, AL045794, H00069,
		AL043423,	AL036924, AL037082, AL038851,
		AL037104,	AL036117, AL036238, T23947, AL036190,
		AL036679,	AW451070, AL036733, Z99396, AW452756,
-		AL037081,	AL037027, AL037601, AL036191, T02921,
		AL037178,	AI535983, AL036158, D51250, AL036765,
		AL036998,	AI535783, AL037054, AL036964, R47228,
		AL036174,	AL037177, AL037021, AL037643, T23659,
		AL037600,	D80253, AL037049, AL037124, AL036858,
		AL037077,	AL036139, AL119457, D59787, AL036132,
		AL036167,	D80043, AL036268, D59275, AL037085,
		AW450376,	
		AL119399,	T48598, AA514190, Z25782, AL036900,
		AL038447,	
		AL042382,	
		AL036227,	-
		AL036719,	AA631969, AL036150, AL037002, D51423,
		T11051, A	AI763414, AL042745, AL119511, AL036999,
-		AL119748,	AI174394, AL040243, AL037679,
		AL042628,	AL037569, D80210, Z25783, AI696819,
_		AW151136,	AL047675, AL079741, AL046356, D59619,
		AW029611,	AI280732, AL045266, AL079977,
		AW071349,	AI608936, AL042744, AI249877,
		AL045620,	AL046926, AI591407, AW089179,
		AL047092,	AL045163, AL039276, H00072, AL121286,
		AI433976,	AI680162, AL045500, AL042787,
		AI433157,	AI554821, AL049085, AI539771,
		AI537677,	AI432666, AI500659, AI815232,
		AI648502,	AI805769, AI801325, AI648663,
		AI500523.	AT625467, AT582932, AT923989.

	, AIS00706,
	62, AI284509,
	, AI433968, AI866573, AI63349
	, AI888661, AI284513,
	, AI633419,
-	03, AL045774,
	20284, AI917963, N80094, AI
	, AW190042,
	, AI699011,
	809, AW151785, AI537515,
	_
	AW193026,
	, AI859464,
_	, AI25183(
	, X68127, AF118808, Z9
	•
	3, AJ244004, AR031375, I18371
	A44171, AR018924, X73004, A63067,
	A51047, A63064, AR018923,
	A48775,
	A38214, A58521, AR015960, I56772, I95540,
	7, AR015961, AR0209
	A25909, I19516, AJ230933, A93963, A93964,
	A95052, A64081, AR043602,
	l, A95117
	A23334, A75888, I7
	2, A23998, I60241,
	AR067731, AR0371
	13189, A43188,
_	3, A81878, I66495
	7, I66496,
	A24782, A35536, A35537, AR022240, A02135,

		-		A02136, A04663, A04664, A11245, A02710, E12615,
		-		3, E14304, A07700, I00074
				A13392, A13393, I19517, A27396, A76773, A22413,
	_			I28266, I21869, I13349, AR027100, A49045,
		-		E03165, E16636, I2692
				I26927, A58525,
				A51384, I03665, Y1
				A70040, AF156294, A97211, E16590, E00523,
				AR038286, I25041, I92483, AR000006, AR038762,
				D88984, I49890, I44516, U87250, A92636, I00079,
				100077, AR008430, AR035975, AR035974, AR035977,
				AF096810, A18722, A91754, AB012117, A97221,
				AF156303, AF156302, X58217, AR064706, I07429,
				I68636, M32676, AF156304, A10361, AF156299,
				A60957, I84554, I84553, A60968, AF096793,
				X15418, I69350, AF130655, AR027069, A10363,
				A52326, A04710, IO
				S83538, Y11447, AR063812,
	•			S
				AR066494, AR060234, I03663, AL137271, A02711,
				AF183393, AL117585, AJ000937, I89947, I48978,
				U80742, AL137463
1977	H2LAS29	887155	Preferably excluded from the	AW408152, AW263155, AA360413, AA314512
			present invention are one or more	
			polynucleotides comprising a	
	_		nucleotide sequence described by	
			the general formula of a-b, where a	
			teger	
			SEQ ID NO:1977, b is an integer of	
			15 to 498, where both a and b	
			d to the positions of	
	3		nucleotide residues shown in SEQ ID	

				E02832, E02493, E04897, E02578, I48917, E02708, E02579 E02709 E01404 E01403 E01405 X51935
				6, E06063, I56011, L03546, X85801, E0085
				E06847, E06846, E03402, I0576
				X01648, X65651,
				647, E01176, A76865, Z36790
				I01586, E01178, E01604, E01177,
				E03605, E03858, A31147,
				A31178, A31148, A07733, A07732, A31150, A31181,
				A31151
1980	HADME31	887280	Preferably excluded from the	AI376391, AW044644, AA435896, AI306612,
			present invention are one or more	AA824370, AA626315, AA991266, AI192974, N78952,
_			polynucleotides comprising a	AI401045, N78829, AI077370, AA448861, W68342,
			nucleotide sequence described by	AA724792, AI708684, AI370929, AI015595,
	٠		the general formula of a-b, where a	AI401211, AW043992, AA862620, AI201717,
			is any integer between 1 to 901 of	AW005929, AI498880, AI718029, AI333236, W93038,
			SEQ ID NO:1980, b is an integer of	AI092949, AI147031, AI004135, W17346, AA027214,
			15 to 915, where both a and b	, T98518, A
			correspond to the positions of	AA878662, W17259, D80253, D80043, D80219,
			nucleotide residues shown in SEQ ID	
	-	_	NO:1980, and where b is greater	D51250, W68383, D80240, AI541365, AA585356,
			than or equal to a + 14.	D80210, D51423, AA585440, AI
				AI546855
				AI546828,
				080196, C142
				D80949, AI557262, T11028, AI536138, D59927,
		_		AI546999, AI525306, AI557238, AI143531,
				AI547039, D80168, AI342055, D80366, AI541205,
				AA585453, AA585434, AI541535, AI541307, T11051,
				AA585476, D57491, AI556967, AI557799, C16300,
				R29445, AI525431, AI546945, AI540967, D81026,
				AI557082, D50995, C14014, AI541534, C16305,
				AI525856, AI525320, AI557808, AI525328,
				AI526194, C75259, AL040155, AL041346, AL041096,
				AL047012, AL041358, AL041277, AL041163,

	AL041098,	AL040621,	AL043538,	AL041324,	
	AL040464,	AL044162,	AL041086,	AL043496,	
	AL041296,	AL041233,	AIS57084,	AIS46875,	
	AI557787,	AL039156,	AL043441,	AL041140, V	W25674,
	AL039150,	AJ239433,	AL038821,	AL039085,	
	AL040193,	AI541013,	AL043445,	AI525653,	
	AI526184,	AA585155,	AI535813,	T24119, T2398	1985,
	AL040149,	AI546899,	AL045725,	AL041197, D61254	61
	AL043612,	T24112, A	.039564, AI	T24112, AL039564, AL039538, AIS	57807,
	AL039108,	AL039678,	AI526196,	AL039915,	
	AL039074,	AL038837,	AL039625,	AL039648,	
	AL039629,	T23888, A	AI541048, AI	AL037726, AL038531	38531,
	AL039109,	AL040992,	AL039924,	AL040463,	
	AL039128,	AL044407,	AL039386,	AL036973,	
	AL045337,	AL037051,	AL039509,	AL045353,	
	AL036725,	AL039423,	AI546891,	AL047219,	
	AL041227,	AL039566,	T23947, AI	AL047057, D59	D59889,
	AL039659,	AL047170,			AL047036,
	AL041292,	D55233, A		AL041051, ALC	AL047183,
	AL040322,	AL041131,	AL046330,	AL045341,	
	AL041133,	AI541509,	AL041238,	AL041142,	
	AL045817,	AL045794,		, AL040529,	
	AL040625,	AL040510,	R29218, AI	AL042909, AL	AL043467,
	AL044186,	AL044037,	AL040091,	AL040128,	
	AL040168,	AL040255,	AL040285,	AL040342,	
	AL040332,	AL040617,	AL045684,	AL040745,	
	AL041347,	AL040370,	AL043677,	AL046442,	-
	AL040553,	AL040839,	AL041752,	AL043444,	
	AL043775,		AL043492,	AL041602,	-
	AR017907,	m		1062873,	34
	A20702, A	A20700, A43		3, A84772, A	A84775,
	A84776, A	A84773, A84		7	32,
	A58522, A	A91750, U87	U87250, A02713	A18053,	A95051,
	I06859, A		A75888, I.70384,	1, A18050, A	A60111,
		AR007512, A	R043601, AS	55	37,
	A35536, A	A02136, A04	A04664, A0213	A02135, A04663, I	E13740,

	A11245, I60241, E12615, A02710, I60242,
	AR035193, A07700, A13393, A13392, A92133,
	AR027100, I66498, I66497, I66496, I28266,
	I66486, I21869, A70040, I84554, I84553, I08051,
	I19525, A25909, A6
	AR025207, Y17188, A85396, I44681, A85477,
	A86792, A44171, AR038855, I66495, I66494,
	M28262, AR066482, I68636, AR035975, AR035977,
	AR031374, AR031375, A85395
	118371, A60985, AR020969, A60990, A85476,
	A91754, A62298, AR037157, AR008430, AF082186,
-	AR035974, AR035976, AR035978, AJ244004,
	ഗ
	D14548, AR038762, I63120, A98767, U94592,
	AR06381
	X83865, I15717, A
	I08396, AF118808, A95117, I08395, X73004,
	AR018924, AR018923, A48774, A48775, AR015960,
	AR000007, AR015961, X55486, I19516, S70644,
	A23998, A95052, AR043602,
	I00074, I92483, AR038286, E03627
	A60209, A60210, A60211, I62368,
	A24783, A24782, I03665, A64081,
	, D50010, A81878,
	A15078, E00523,
	E00609, A64973, I49890, A11178, E01007,
	903, D285
	A22413, E16590, A4
	E16636, Z32836, AF156294,
	2, AF149828, I25027, I26929, I4451
	T26927 T25041 AR031488

				113521, I52048, I44531, A90655, X58217, Y11923,
				144516, AR031566, A58525, I01995
				, AJ230933, AF019720, I18895, E03165,
				:0699, E00696, E00697
1981	HFVJL45	887399	Preferably excluded from the	AI074616, AW008223, AI523733
			present invention are one or more	AI309184, AI910363,
			polynucleotides comprising a	AI266526,
			nucleotide sequence described by	, AA860930
			the general formula of a-b, where a	R95740, AA256366, R95884, AW449536, AI027719,
			is any integer between 1 to 1413 of	H80516, AI674127, AI202271, T57140, T11308,
			SEQ ID NO:1981, b is an integer of	AI247797,
			correspond to the positions of	œ
			nucleotide residues shown in SEQ ID	
				AW384394,
			a	AT.119439
			3	
•				AL119496, U46350, AL134518, U46349, AL119444,
				U46347, U46351, U46341, AI142132, Z99396,
				AL119355, AL119483, AL042614, AL119396, U46345,
				AL134538, AI142137, AL134530, AL134519,
				AL134531, AL119401, AL079687, AL037205,
				AL042980, AL042896, AL043037, L48516, AC004022,
				L76193, AC005021, AB026436, AR060234, A81671,
				AR054110, AR066494, AR069079
1982	HWLFE56	887421	Preferably excluded from the	AF061056, AF084644, AF084645, AJ009937, AJ009936
			present invention are one or more	
			polynucleotides comprising a	
			nucleotide sequence described by	
			the general formula of a-b, where a	
			is any integer between 1 to 697 of	
	-		SEQ ID NO:1982, b is an integer of	
			15 to 711, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1982, and where b is greater	

			than or equal to a + 14.	
1983	HSWBP93	887475	Preferably excluded from the	
			present invention are one or more	AA974489, AA249308
			polynucleotides comprising a	
			nucleotide sequence described by	
			al formula of a-b, whe:	
			is any integer between 1 to 509 of	
			SEQ ID NO:1983, b is an integer of	
			15 to 523, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1983, and where b is greater	
			than or equal to a + 14.	
1984	HSLJF91	887535	Preferably excluded from the	AI525881, D78870, H11172, R19956, AA308077,
			present invention are one or more	AI591060, AA350839, AI557291, AF091352, A64392,
			polynucleotides comprising a	AB021221, S82167, X62568, M32977, A64394,
			nucleotide sequence described by	A64398, A64402, AF022375, A92244, A64400,
			the general formula of a-b, where a	X81380, M31836, M32976, AF071015, AF133248,
			is any integer between 1 to 450 of	A92248, S85192, AJ010438, A92246, M27281,
			SEQ ID NO:1984, b is an integer of	A64396, A92242, AF214570, E13215, AF186236,
			15 to 464, where both a and b	E15157, M32167, M33750, S38083, X89506,
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	AF222779, AF215725, L20913, S38100, S37052,
			NO:1984, and where b is greater	AF062645, AF106942, AF022179, S85199
			than or equal to a + 14.	
1985	HKLSC61	887803	Preferably excluded from the	AL039924, AL045794, AW013814, T02921, T24119,
			present invention are one or more	
			polynucleotides comprising a	D59787, D80219, AL039629, AL039625, AL039648,
			nucleotide sequence described by	AL038837, AL039074, AL037726, AL039678,
			the general formula of a-b, where a	AL039108, AL039538, AL039564, AL039156, D59275,
			is any integer between 1 to 1219 of	AL039659, AL039566, AL039509, AL039150, D80227,
			SEQ ID NO:1985, b is an integer of	AL044530, AL038531, AL039109, AL038821,
			15 to 1233, where both a and b	AL040992, H00069, AL043423, AL039128, AL044407,
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	AL039386, AL039476, AL045341, AL039423,
			NO:1985, and where b is greater	AL042909, AL043441, AL044412, AL039410,

	than or equal to a + 14.	AL044364, AL043445, AL038025, AL043422, D80210,
		AL036725, D51423, D80134, D59619, D80391,
		D8
		D80949, C14227, AW450335, AL039521, AL039085,
		AL036196, T23947, AL037526, AI535783, AL037639,
		AW451070, D80366, D80168, AL037615, D80045,
		3, AW452756, T11051, D81026,
		L036117, AL
		1,
		AL036238, C15076, AL036733, AL037082, AL036679,
		AL038851, D80022, AL036418, D80038, T23659,
		AL037054, D80195, AL037027, AL036765, AL039504,
		AL036158, D58283, T11417, D81030, C14429,
		AW293068, D80188, AL037047, AL036964, AL036190,
		D51799, D80378, D59467, AL036650, F13647,
		AL036191, AL037104, T03269, AL037177, AL036998,
		AL037679, D50979, D80522, T48598, D80212,
		C14298,
		Z21582, AL036132, AL036167,
		AL037600, AW450376, AA514190, D80164, C14331,
		D59695, D80166, AI021934, AL
		W206560, D80268, AL036152,
		2, AL037021, D52291,
	-	Z99396, AL036900, AL036
		AL044447, D59610, AL037085,
-		I910186, D81111, C14407, C14389,
		AW451416, H00072, T23656, AL037081, D51060,
		AL036228, AW178893, AA305409, AL037077,
		AW178775, D80014, AW378532, D80248, AL036808,
		v
		.036858, AW
		1, AL037002, AW360
	-	6, AW360811, AI557774, AW3
	-	AW352117, T02974, Z25783, AL039417, C05695,

		AL044413,
		AW360844,
		AW360817, AW375406, AW378534, AW377672, D80439,
		, A85396, A25909,
		A44171, AR062871, A84775,
		AR017907, A84774, AR067731, AR067732, A20702,
		A58522, A91750, A43189, A43188, A20700,
		A38214, A95117, A95052, I56772, I95540,
		, AR031374, A93963, A939
		A18053, A51047, A63064, AR018923, A49700,
		I18371, A48774, AR031375, AR043602, A63072,
		m
		A75888, I70384, AR068506, A18050, A60111,
		A23633, AR015960, A23998, AR007512, AR000007,
		AR015961, A58521, I63120, I60241, I60242,
		AR054109
		A58524, E12615, AR035193, A92133, A24783,
		2, A58523, E14304, A27396, AR02710
		.9045, E16678, A8
	*	, A93016
		I26928, I26930, I26927, A58525, E13740, I49890,
		vo
		U87250, A13038, A29289, AJ244004, D34614,
		AR008430, AR029417, A71435, AB012117, I13349,
		Z96142, A97211, A07699, E08322, I74623, A71440,
		V00745, AF156303, Y17188, A02712, X73004,
), AR028668, AR0286
		AR028670, A68112, A68104, AR067733, AF118808,
		AR029418, I52048, I44531, AR067734, A84746,
		AR028672, I66498, I66497, I66496, AR038066,
:		I50882, I66486, I66487, I15353, I19516, A83643,

	AA989485, T98916, AA487702, AA297484, AA297153,
	AA464649, AA292774, AW170481, AI963760, W79558,
	AA394263, AI986058, AA903542, AW079683,
	\mathbf{H}
	AA287210, AA297527, AA341051, AA861541,
	AA297453, AA557937, AA133595, AA464548,
	AI275661, AI719497,
	AA496440, AA481372, AI673125, AA565649, C21003,
	4298491, AI
	AI273816,
	AA133686, AA327635, AA411355, AJ011497,
-	AL137480, I48978, AF159615, A70386, AF102578,
	A77033, A77035, A08910, A08909, ALO50024,
	AF026816, A08913, Z37987, U73682, AL110280,
	AL117435, AL080159, AL035458, AL122050,
	AF087943, AL117457, AR068753, AL137533,
	AR034821, AF113019, A07588, S36676, S83440,
	X82434,
	Z82022, I25049, AF185576, AL080126, AF057300,
	I89931, S63521, A65341, AF0
	AJ005690, I49625, AF119336, AJ00
	, AF026124
	U49908, AL117635,
	AL110296, I66342, X83508, Al
	AF067728, X80340,
	AL050138, AF106862
	3, S61953, AL137283,
	AL133619, AL137521, X72889, AL137478, AL137560,

				AF079763, AL110221, I25048, AF162270, AL117648,
				L13297, AR000496, A93914, U39656, AF090900,
				I09499, AF182215, AL133560, AB031064, AR020905,
				AL133637, U92992, AF100931, X66862, AF054599,
				AL049938, AL133557, A93350, AL096744, AL050146,
•				AF061981, A52563, X66366, AJ012755, AL080118,
				X61970, U75932, AF113694, E03348, AL133080,
				AF051325, U58996, X84990, E01314, AF118558,
				AL049452, AL133031, AF061573, AF124435,
				AF076464, AL050277, A65340, AL049283, I33391,
				M30514, L3
				AL137558, AL117440, AL049447, AL133067,
				AL133084, AL137557, AF118070, AL133640,
				AL117626, 126207, AF111851, A45787, AF106657,
•				X98834, AF017437, A08908, AL049300, AL080146,
-				
-				
				1000001日本
				AF162/82
1987	HWLOA40	887892	Preferably excluded from the	AA297147,
			present invention are one or more	AW300770, AI691072, AA563933, I95745
			polynucleotides comprising a	
			nucleotide sequence described by	
			the general formula of a-b, where a	
•			is any integer between 1 to 507 of	
-			SEQ ID NO:1987, b is an integer of	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1987, and where b is greater	
			than or equal to a + 14.	
1988	HCQCF10	887936	Preferably excluded from the	AI82360
			present invention are one or more	AA410501, H66313, W37614, AF131758
			polynucleotides comprising a	
			nucleotide sequence described by	
	T		l	

			the general formula of a-b, where a	
•			iny integer between	-
			SEQ ID NO:1988, b is an integer of	
			15 to 346, where both a and b	•
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:1988, and where b is greater	
			than or equal to a + 14.	
1989	HAIBW90	887996	Preferably excluded from the	AA481723, AA626700,
			present invention are one or more	AA47933
-	•		polynucleotides comprising a	AA479335, AA165042,
			nucleotide sequence described by	AI400160, AW370132, AI924188, AW015034, F06368,
			the general formula of a-b, where a	C15288, H89161, AA364967, AW262875, AI566873,
			is any integer between 1 to 938 of	AA371283, AI566669, AI864174, AA304171,
			SEQ ID NO:1989, b is an integer of	
			15 to 952, where both a and b	~
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	AA934901, N40173, R46865, AW157527, AI374781,
			NO:1989, and where b is greater	AI379523, H64413, AI371781, R78607, AW173107,
				, AA195689,
				AI742505, AI087379, AA527113, AA527036,
				AA373921, AI952545, AI269215, AI245243,
•				AA302499, AI792601, AA600140, AI040546, H92421,
				C16267, AI805770, Z24901, AA625963, AI139790,
				AI360032, N40209, AI084568, D57610, AI753737,
	. •			C16455, R35721, AA159931, AI024890, AI869836,
	•			AI829158, AI804015, AA477326, AA430365,
				AI640196, N30689, AI371005, AA478600, AA256968,
				AA021044, AA657967, AW072764, N41298, AA905154,
				AA758776, AI955815, AA865424, AI857650,
				AI091988, AW242058, H92638, AA234867, AI864141,
				AA252106, AA424350, AA302462, AI468749,
				AW090440, AI336687, AA732498, AA302463, C16184,
				608, AW085952, AI934133, AI2695
				F05286, R46768, C16334, AB006077, AF006484

07070, D80268, D80366, F13647, 015, D80522, AA305578, C14227, 995, AW177440, D51022, D81026, 111, D80391, D80248, D59787, P 283, D59619, D80210, D80240, C	AA514188, D80195, AA305409, D80196, D59859, D80022, D80043, D80166, D59927, D59467, D51423, D51799, D80164, D59275, D80253, D80038, D80227, D59502, D80212, D81030, D80219, D51060, D80188.	, AA514186; D59889, D80439, C15076, D80269, D59610, D57483, D80193, D8004, T03116, D80247, D80064, AW378533,	D80378, D51759, D80241, C14014, 103269, AW178893, D80133, AW178906, D80302, D80168, C14407, D80157, AW360811, AW178759, D51103,	AW378540, D80251, D80949, AW35212	366296, AW360	T02974, AW179020, AW375406, T48593, AW378534, AW179332, AW377672, AW378528, AW179023,	AW178905, AW352158, D51250, AW17731, AW178762, AW178754, AW179024, D59373, C05763,	IS57751, D80134, H67854, C03092,	H67866, D80132, AA809122, AW179004, AW360834,	0, AW178907, AW178908, AI5259), AW378520,	D3931/, D38246, D80238, AW1/9012, AW1/8980, D80014, AW177733, D59503, AW179018, AW178914,	4, C14046,	4, D58101,	6, AW378543	1, D59551, AI52527, D80228, AA285331,	8, AI525235, C16955, AI525912, AI525	905856, D45273, A1525242, 233452, A1525 525237, A1525215, AA305720, AW378542, C
more d by	leger between 1 to 592 of 1990, b is an integer of where both a and b	e residues shown in SEQ ID and where b is greater qual to a + 14.	Q & U	UA	. 4	H	4 4	Α	H.F.		A (4			at ·	4.6	A
888041																		
H2CBE03												•						
0661																		-

			H67858, T03048, AI525222, T02868, F13796, AM3460855, Z30160, D31458, D51053, D79997.
			, A84916,
			A62298, AR018138, AR008278, AF058696, AB028859,
			5, AR060385
			IS0128, IS0133, X67155,
			39785, A78862, D346
			4, X68127, A94995, AR060138,
			AR052274, AR066488, Y09669, A43192,
-			A43190, AR038669, I14842, AR008443, AR066487,
			17187, A63261,
			X82626, AR008277, AR008281, D50010, AR062872,
			Å70867, AR016691, AR016690, U46128, AR016808,
			AR008408, AR025207, X64588, A64136, A68321,
			I79511, D13509, AR060133, I18367, AF123263
1991 HE90119	888051	Preferably excluded from the	AL043100, AL045367, AL042404, AA326785, R34387,
,		present invention are one or more	
		polynucleotides comprising a	U82536, AF097999, AF098010, AF098011, AL050372
		nucleotide sequence described by	
		the general formula of a-b, where a	
		is any integer between 1 to 1083 of	
	100	SEQ ID NO:1991, b is an integer of	
		15 to 1097, where both a and b	
		correspond to the positions of	
		nucleotide residues shown in SEQ ID	
		NO:1991, and where b is greater	
		than or equal to a + 14.	
1992 HJACE25	888063	Preferably excluded from the	AA311008,
		present invention are one or more	_
		otides comp	, N30086,
		nucleotide sequence described by	AI918715, D80391, D80196, AI282428, D59787,
		the general formula of a-b, where a	
		is any integer between 1 to 889 of	, D80253, D59619, D80210, D80240,
		:1992, b is an inte	, D80212, D81030, D57483, D80195, D5988
		15 to 903, where both a and b	9, D59610, D80043, D59467, D59502, D5992
		correspond to the positions of	D80022, D80366, D59275, D80193, D80241, D80378,

	nucleotide residues shown in SEO ID	D80024, D50995, D50979, C75259, C14429, D80164,
	and where b is great	3269, D80045,
	equal to a + 14.	D51060, AA305409, AA352266, D80134, AW178893,
		D51250, C14227, D81026, D80949, D80268, F13647,
-		
-		D51022, D80522, D81111, AW179328, Z21582,
		8, AW378532, AA305578, D59695,
		D80251, AW177511, AW369651, AA557885, D52291,
		D80064, AA514186, D80133, AW352117, D51097,
		AA285331, AW360811, AW378540, AW377671, C14407,
		AW375405, AW360844, AW360834, AW366296, D80439,
		D80132, AW360817, AW375406, AW378534, AW352171,
		, AW377672,
		AW178905, AW178754, AW179018, AW179024,
		AW177505,
		1178909, AW
		AW177731, AW178907, AW179019, AW178971, D80247,
		AI557751, AW179004, AW179329, T02974, AW352174,
		, D80014, AW17773
		, T11417, D80157,
		, D51103, D51759, AW
		AW178911, AW352163, D58101
		AI557774, T48593, AW378539, D80258, D59503,
		D51213, D59627, D45260, H67854, D50981,
		950, AW178986, AI525923,
		AA809122, AW177734
		D59474, AI525920, D51221,
		C14344, C14973, AA514184, T03048, AA033512,
		AI525227, AI535686, AW179013, AW178759, D59551,
		AF080255, AF073771, A62298, A84916, A62300,
	-	AJ132110, X67155, AR018138, D89785, Y17188,
		78862, D26022, A25909, D34614
		AR008278, X82626,
		AB012117, Y12724, X68127, A85396, AR066482,

				A44171, A8	A85477, I19525, A86792, A82595, U87250,
				AR008443,	AKUGU303, ABUU2443, AKU1900 6, IS0132, IS0128, IS0133,
_				AR066488,	AR066488, AF135125, AR016514, AR060138, A45456,
				A26615, AR	A26615, AR052274, Y09669, A43192, A43190,
				AR038669,	4.
					I14842, AR054175, AR008277, AR008281,
				_	Y17187, X64588, AB033111, A63261,
		-		AR064240,	AR008408, AR062872, A70867, AR016691,
	_			AR016690,	U46128, D13509, A64136, A68321,
				~	
				AF123263,	X93535, AR008382
1993	HMWIR85	888153	Preferably excluded from the	AA195033,	AW150723, AI805372, AI826894,
			present invention are one or more	AW245532,	AW250255, AW269478, AI929681,
			polynucleotides comprising a	AI814415,	AI984552, AI081263, AW178616,
			nucleotide sequence described by	AW352048,	AW352014, AW250589, AW178530,
			the general formula of a-b, where a	AI688093,	AW352019, AW178493, AW178640,
			is any integer between 1 to 2985 of	AA861507,	AW178500, AI146435, AW178537,
			SEQ ID NO:1993, b is an integer of	AA514698,	AW352042, AW352039,
		_	15 to 2999, where both a and b	AW352051,	AW178535, AW
			correspond to the positions of	AW178504,	AW352041, AW352035, AA936386,
			nucleotide residues shown in SEQ ID	AA573323,	AW178641, AW178529, AI566475,
			NO:1993, and where b is greater	AA928767,	AI963685, AW178642, AA826410,
			than or equal to a + 14.	AW178605,	AA648798, AA250731, AW178506,
				AI360338,	AA865431, AI342420, AI439684,
				AI351346,	, AI355698,
				AI308956,	, AW178495,
				AI015535,	AI096589, AI683046, AI884370,
				AI473866,	AW178634, AW178667, AA495743,
				AW178531,	AW178614, AA196630, AA533557,
				AA122301,	AI090332, N53164, AA024938, AW178637,
				AI370758,	AW178536, AW366100, AI832020,
				AI859889,	AI571925, AI274028, AA024855,
				AA206040,	AA583100, AW178507, AW178533,
				AW178615,	AI289830, AW352018, AA636082,
				AW178672,	AI220039, AA654736, AI831555,

				AW246444, AI080500, AW352027, AW366090,
				AW178511, AI289407, AA478574, AW366108,
	•			AW366085, AW178670, N93845, AA640678, N35617,
				AW366113, AA665800, AI971078, AW352022,
				AA482749, AW366082, T35187, AA187140, AA732528,
	_			AL040485, AI699027, AL048191, AL048192,
				, AI446512, AW366088,
				AA478709, AW178516, AA759075, AW366107,
				AA313616, AA122275, AW178503, AI970947,
				AW130860, AI051515, AW366098, AA651674,
•				AA211795, AA211028, AW366081, AW366095, R56468,
_				H72835, AI300727, N58601, R56467, AA471174,
				AW366092, AA074578, AA173306, AI867698, R43285,
				AA494312, AW178538, F089
				, AA636095
	-			AW352028, W44497, F11265, AI929035, AA828212,
-				, AA456024,
				AW178527, AA181660, T3
				533, AW178630, AI2501
				, Z28803, AW1786
				T74206, H78935, H80408, T35534, T74205, F15248,
				471318, H71931, AW352024, AW352025,
				7, AI915730, AA214391, W26
				1061, AW351670,
				AW178526, T35316, F15255, R18867, AW352033,
				AI274653, AA092564, R43478, AA480524, R37293,
				AW178494, AA404508,
_				AI587130, AW178674, W27395, T81825, AA258411,
				AW178518, AW366099,
				AA903360, AF046001, AC005899, AB013357, X74802,
				8362
				52
1994 H	HCRPV38	888254	Preferably excluded from the	W68102, AA005326, AA447946, AA101751, W67683,

present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 324 of 15 to 338, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:1994, and where b is greater than or equal to a + 14.	Preferably excluded from the A1992179, AM188159, A1926499, A1926498, present invention are one or more A1869696, H38016, A4811687, AA301183, A1018137, A1869689, AA974505, A1090091, AA452882, nucleotide sequence described by A4866789, AA974505, A1090091, AA452882, A4866789, AA974505, A1090091, AA452882, A4866789, AA974605, A1090091, AA452882, A4918138, AA746089, AA974498, N21230, AA905692, SEQ ID NO:1995, b is an integer of A1123002, A1223636, A1361685, AA431160, A1123002, A1223636, A1361685, AA431160, A1123002, A122440, N27905, AA431160, A112302, A112304, A1125440, N27905, A1239567, A1720492, NO:1995, and where b is greater A1467630, W70189, A1125441, AA486664, AA047262, A446630, AA965328, AA6632987, AA761345, A4467630, W70189, A1125440, A279411, AA486664, A37709, A4467630, AA963525, AA622987, AA761345, A4467630, AA963525, AA622987, AA76936, A4467630, AA963525, AA632987, AA76936, A4467630, AA963525, AA95388, N21070, AA884028, AA846912, WA2764, M27315, W22674, A1381262, R51770, A4467630, AA963825, AA358806, AA883739, CA2686, AA883739, A496864, A112484, AA963868, T81857, AA88188, A496864, A112484, AA96886, T81857, AA88188, A496864, A118484, AA96886, T81857, AA88188, A496864, A118484, AA96886, T81857, AA88189, A496864, A118484, AA96886, T81857, AA88189, A496864, A118484, AA96886, T81857, AA88189, A496864, A1184844, AA96886, AA8888, AA88888,
present invention arrepolynucleotides componucleotide sequence the general formula is any integer betwee SEQ ID NO:1994, b is 15 to 338, where bot correspond to the ponucleotide residues NO:1994, and where b than or equal to a +	Preferably excluded present invention ary polynucleotides compounded sequence the general formula is any integer betwee SEQ ID NO:1995, b is 15 to 2346, where bo correspond to the ponucleotide residues NO:1995, and where by than or equal to a +
	888402
	HSRBB92
	1995 HSR

				AA707548,	AW340816,	AA613385, D61871,	D61871, H3824	242,
				H81487, A	AI218047, AI190091, AA453052,	1190091, AZ		AW138451,
				AW294322,	AW452108,	AI766143,	æ	
				AI832222,	AW292106,	AJ271408,	AF132938,	
				AF106798,	AL133631,	AR007449,	U39643, AF0	AF094700
9661	HSYEA10	888523	Preferably excluded from the	AI037890,	AW003999,	AI858060,	AW084608,	
			present invention are one or more	AI589010,	AW304188,	AW117854,	AI038497,	
		he la terrati	polynucleotides comprising a	AI452673,	AI743739,	AI147810,	AA181048,	
			nucleotide sequence described by	AA187507,	AA081006,	AA082736,	AA187264, N	N94407,
			the general formula of a-b, where a	AA187361,	AA181882,	AI079886,	AA181880,	
			is any integer between 1 to 2007 of	AA188249,	AI445147,	AI471432,		AA100829,
			SEQ ID NO:1996, b is an integer of	AA503656,	AA081230,	AA182826,	W47343, AA1	AA182830,
			15 to 2021, where both a and b	AA181134,	AI085755,	AA132297,	AI076956,	
			correspond to the positions of	AI613182,	AA081149,	AA188049,	AA186634,	
			nucleotide residues shown in SEQ ID	AI081490,	AA186808,	AI918426,	AA186376,	
			NO:1996, and where b is greater	AA081282,	AA082516,	AA186389,	AA081208,	
			equal to a + 14.	AA582862,	AA147528,	AA157628,	AI082493,	
				AI282835,	N94510, W4	19497, AA1E	N94510, W49497, AA181875, AA191501,	501,
				AA083542,	AA157752,	W47445, AA101069,	1101069, AA1	AA186754,
				AA081283,	AA182682,		C06085, W39354	354,
				AI800644,	AA157468,	AA186973, AA374217,	AA374217,	
				AA386155,	W23960, T2	27821, AA08	T27821, AA083575, AI654536	536,
				AA308204,	W52714, A	4852603, A	AA852603, AI270203, AA188296	88296,
		in		AA852324,	AA852602,	AA143331,	AW449628,	
				AA083541,	AA372360,	AA158121,	AA186524,	
				AA304334,	AI932880,	AA187348,	W60270, AA3	AA308786,
				AA188042,	AA157416,	T18504, AA143201,	1143201, C02231	231,
		·		C02091, A	C02091, AA156273, AA157642, AA100067,	A157642, A	A100067, W56826	826,
				W56827, A	A514656, AV	4376428, W	AA514656, AW376428, W31070, AW37642	420,
				AI912469,	X54925, X(35231, IOIC	AI912469, X54925, X05231, I01070, AF14888:	2,
				X54724, X	X58256, A47086, U78045, S75623,	386, U78045	5, S75623, M	M17821,
				M15996, M	M17822, M17823, M16567, U78629	323, M16567	7, U78629,	-
				AJ002550,	M25663, AF	AR040773, AF	AF023338	
1997	HE2CC22	888673	Preferably excluded from the	AW368993,	AI638166,		AI041204,	
			present invention are one or more	AL042348,	A1478737,	AI760185,	AI830441,	
			polynucleotides comprising a	AI126299,	AI217176,	AI092924,	AI799277,	

			nucleotide sequence described by	AI857759,	AA993596,	AI381442, AI62034	345,
			äl	AI027099,	AA743334,	AI827435, AI13880	305,
			is any integer between 1 to 1941 of	AA136171,	AI285950,	AI635387, AA664373	373,
			SEQ ID NO:1997, b is an integer of	AI827427,	AI015864,	AI222122, AA843185	185,
			15 to 1955, where both a and b	AA976953,	AW021642,	AI685358, AW195005	005,
			correspond to the positions of	AI206601,	AW023027,	AW450169, R80985,	
			nucleotide residues shown in SEQ ID	T78995, A	4912496, A	T78995, AA912496, AA926963, AW451943,	3, AI249890,
			NO:1997, and where b is greater	AW269181,	AW026792,	R68431, AA731014,	1, AW074050,
			than or equal to a + 14.	AA922059,	AA757551,	H12605, AA689507,	7, W79832,
				AA412149,	AW135157,	AW071659, R49066,	s, AA056573,
				AA278795,	H91438, AI567760,	H12655,	
				AA040923,	AA721747,	T78939, Z41658,	AI767505,
				AA766306,	AA987389,	AI538809, R68430,), R26542,
				AA056678,	AA353814,	H91332, R80785, R25352,	R25352,
				AA361014,	AA536104,	AI699602, R57916,	5, AA278600,
				AA040922,	AB007949,	X65024, D21089	
1998	HOUAC22	888708	Preferably excluded from the	AI821479,	AI739517,	AW082828, AA533173	173,
			present invention are one or more	AI198451,	AA532999,	AI821509, AI791624,	524, U25936,
			polynucleotides comprising a	AA315607,	AI000331,	AW139172, AA35887	375,
			nucleotide sequence described by	AI125295,	AI216275,	AW005074	
			the general formula of a-b, where a				
			is any integer between 1 to 1144 of				
			SEQ ID NO:1998, b is an integer of				
			15 to 1158, where both a and b				
			correspond to the positions of				
			$\boldsymbol{\sigma}$				
			, and where				
			than or equal to a + 14.				
1999	HHECU01	888720	\Box	AA853396,	AC005041		
			present invention are one or more				
			polynucleotides comprising a				
			nucleotide sequence described by				
			the general formula of a-b, where a				
			is any integer between 1 to 1113 of				
			SEQ ID NO:1999, b is an integer of				
			15 to 1127, where both a and b				

			correspond to the positions of					
			nucleotide residues shown in SEQ ID					
			NO:1999, and where b is greater					
			than or equal to a + 14.					
2000	H2LAP34	888783	Preferably excluded from the	AA314278,	AA315476, 1	AA133008,	AW301013,	
			present invention are one or more	AA314092,	AA386092, AA411572, AA427682,	AA411572,	AA427682,	
			polynucleotides comprising a	AA315987,	U46281, W76038,		W42816, AA314613,	513,
			nucleotide sequence described by	AA477668,	H52355, C1	C17482, AA47	AA477851, AA4813	31359,
			the general formula of a-b, where a	R83104, A	R83104, AA410758, W02292,		W79944, AA329443,	143,
			is any integer between 1 to 464 of	R46315, W	W07627, AW366382,		AA335138, R83126,	
			SEQ ID NO:2000, b is an integer of	AA302305,	W19402, H27934,		AA659027, AA411998,	11998,
			15 to 478, where both a and b	AA151635,	AA366470, AA358810, AA053648, T49358	AA358810,	AA053648,	T49358,
		•	correspond to the positions of	AA378171,	R48529, AA159070, AA838273, T62103,	159070, AA	.838273, Te	52103,
			nucleotide residues shown in SEQ ID	AA429117,	AA158752, AA134180, AW37622	AA134180,	AW376226,	
			NO:2000, and where b is greater	AA149262,	AA410673, 1	U92985, AR065358	065358	
			•					
2001	HNTAR08	888950		AW236102,	AA218985,	AA906740,	AA737950,	
			present invention are one or more	AA220991,	AA926805,	AA206111,	AA206112,	
			polynucleotides comprising a	AI653195,	AA865714,	AA220997,	AA968722,	
			nucleotide sequence described by	AA218991,		AI357043,	AI652879,	
			the general formula of a-b, where a	AI970161,	AW025944,	AA902285,	AI655507,	
			is any integer between 1 to 1247 of	AW003483,	AA902779,	AI824839,	AI917697,	-
			SEQ ID NO:2001, b is an integer of	AI671508,	AI962316,	AA074560,	AR040708,	S52658,
			15 to 1261, where both a and b	AR040709				
			correspond to the positions of					
			nucleotide residues shown in SEQ ID					
			NO:2001, and where b is greater					
			than or equal to a + 14.					
2002	HWLWH6	889136	Preferably excluded from the	AI694583,	AA280341,	AW369780,	AI572844,	
	9		present invention are one or more	AA968512,	AI250884, AI798375, AI370669	AI798375,	AI370669,	
			polynucleotides comprising a	AW181892,	T06923, AW293265, AA947819, AA598509	293265, AA	.947819, AJ	4598509,
			nucleotide sequence described by	AL035420,	AL050030, AL022727, AC004129, AC005082	AL022727,	AC004129,	AC005082
			the general formula of a-b, where a					
			is any integer between 1 to 1517 of					
			SEQ ID NO:2002, b is an integer of					
			15 to 1531, where both a and b					

			correspond to the positions of					
		<u>-</u>	de residues shown in					
			002, and where b					
			equal to a	į				
2003	HWLCJ12	889263	\Box	AI632964,	AA826324,	C06338, A]		AA622862,
			present invention are one or more	AI890787,	AA775044,	AA621523,	AA585439,	228355,
			polynucleotides comprising a	AI525556,	AIS41374,	AA585453,	AI535639,	230131,
		_	nucleotide sequence described by	AI546999,	AI546855,	AI541514,	AI525316,	
			formula of a-b, where	AI541510,	AI525306,	AA585101,	AI541523,	
			is any integer between 1 to 2319 of	AI557731,	AA585434,	AIS41534,	AI541365,	
			SEQ ID NO:2003, b is an integer of	AI526140,	AI541509,	AIS46828,	AI525431,	
				AA585440,	AIS56967,	AI526194,	AI541017,	C15189,
				AI540967,	AI547039,	AI557262,	T11028, A	AIS57807,
			nucleotide residues shown in SEQ ID	AI541535,	C16300, A.	C16300, AI557799, AI541205,		AI546945,
			NO:2003, and where b is greater	D61254, R	R29445, AI541307, AI535813, AI557787,	41307, AIS:	35813, AIS	57787,
			equal to a + 14.	AI546899,	AI557238,	R28735, AL040510,	L040510, AI	AL040625,
				AL045817,	AL041142,	AL041238,	AL041133,	
				AL047183,	AL040322,	AL041131,	AL046330,	
				AL041051,	AL041292,	AL040119,	AL047036,	
				AL047170,	AL047057,	AL047219,	AL041227,	
				AI525653,	AL040463,	AL039915,	AL043612,	
				AL041197,	AL040155,	AL041346,	AL040529,	
				AL041096,	AL047012,	AL041358,	AL041277,	
				AL041163,	AL041098,	AL040621,	AL043538,	
				AL041324,	AL040464,	AL044162,	AL041086,	
				AL043496,	AL041296,	AL041233,	AI526180,	
				AL043467,	AL041159,	AL045725,	AL044186,	
				AL041140,	AL040193,	AI557082,	AI526196,	
				AL044037,	AL040091,	AL040128,	AL040168,	
				AL040255,	AL040285,	AL040342,	AL040332,	
				AL040617,	AL040553,	AL045684,	AL040745,	
				AL040370,	AL043677,	AL046442,	AL040839,	
				AL041752,	AL040149,	AL043775,	AL044165,	
				AL043492,	AL041602,	AL045920,	AL041278,	
				AL038838,	AL040253,	AL044074,	AL041635,	
				AL045990,	AL040458,	AL044199,	AL044187,	

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	AL043848, AL041459, AL043570, AL041577,
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	AI541508, AL045671, AL046327, AJ239433,
	AI535660, AL040238, AL041955, AL041347,
	AL038761, AI541048, AL040075, AA585476,
-	, AI540920, T23957, AI526184,
	AI541013,
	', R29177, AI526073, T2
	AL042096, AL037436,
	AI526187, AL039643,
	AL044125, AI557279, D55233, AI541390, AA174170,
	AL045211, AI541356, AI525321,
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	5, E00696, AR05165
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	A70869, E12584, AR035974, AF
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	AR009152, AR050070, X82786, AJ231011, U87250,
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	.8, A06419, A21892, A23997
	A68114, A89633, A89634, A21895, A05160, A08030,

				A20502, X87559, I05488, I61310, A60961, A60977,
				AR002333, A60985
2004	HNGEF72	889299	Preferably excluded from the	
		_	present invention are one or more	
			polynucleotides comprising a	AI806055, R71461, AA306731, AA034255, H53686,
			nucleotide sequence described by	AI741660, H82553, N28450, AI452969, AA318128,
			the general formula of a-b, where a	
			is any integer between 1 to 2385 of	AA025373,
			SEO ID NO:2004, b is an integer of	A156782, U2
				AA303132, AI638569, AI052532, AA091675, R99679,
			correspond to the positions of	AI278003, AI720617, AW051583, AA804776,
			nucleotide residues shown in SEQ ID	AA319103, AW148694, AA029525, AW247858,
			NO:2004, and where b is greater	AW021737, AI140193, AW055259, AA565273,
			Q.	AA642437, AI240825, AI248594, H72148, AA156851,
				AA573394, AA029460, AA359482, T50440, AA018596,
				AA214611, AA634569, AA725707, AA709248,
				AA536183, AW082332, AA361479, AA447253,
				AA447268, AA353770, AI567232, AA962385,
				AA709244, AA767996, AI766591, AI358947, C18192,
				C16865, AW193910, AW235731, AA707012, AW304793,
				;, AI939507, R10615,
				R86259, AI276029, AI561192, N74387, AA131938,
				N74439, AW439563, AA013432, AI753280, AI267829,
		_		AI189108, W04994, R28492, N52383, T85708,
				AL031769, AC007970, AL034426, AC005697,
				AC002065, AC008082, AC006010, AC009286, D87009,
				AC009241, A90827, Z92545, AC009399, AJ243211,
				AL022400, AC004460, AL021917, AC004382,
				AC006522, AC007270, Z99569, AC005323, AC006083,
				AC005681, AC007788, Z98751, AC005731, AC000085,
				AL031684, AC009396, Z97180, U40455, AL117351,
				AC006024, AL109954, AC009514, AC006500,
				AL132994, Z98172, AC002094, AC004553, AC004993,
				88, AL034347, AC003681
				AL109654, U66083, AF109718, AC004844, AL031672,

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 4, AC004875, AL133162, AL133216	
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AC005228,	
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AP000352,	
 AL080317, AL132800, AC004385, Z84470, AC00244	3002449,
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AF205588, Z96050, AC007878, AL049766, AC005326	2005326,
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AL009028, Z72522, AC005145, AC004551, ALI	AL133249,
, AC018633,	U85198,
, AC004474, AE000661, AC007751	
AC004095, AC006036, AC009181, Z98753, AC00	AC006968,

				AL031116.	AL109748,	AP000078,	AC007455,	
				AL031586,		AL022574,	7	
				AL031393,	AC004452,	Z99497, AL137624,		AL079342,
				AB020871,	AC006463,	AC006984,	AC006167,	
				AC004389,	AC004915,	AB023050,	AP000511,	296774,
+	0 2000	0000	brofornhiv evoluded from the	A1952777	AT346020.	AW024883,	AL046029,	
C007	o 	200	y excluded from	AI590661,	ഗ	AW073186,		
			leotides compri	AL037668,	AW151753,	AI419538,	AA399154,	
	-		nucleotide sequence described by	AI420960,	AA971504,	AI424070,	AI983928,	
			the general formula of a-b, where a	AI858710,	AW264165,	AI970601,	AI422333,	
			is any integer between 1 to 1902 of	AA610484,	AA481014,	AA758319,	AA486535,	
		-	SEO ID NO:2005, b is an integer of	AI273879,	AA865664,	AA528037,	AW440638,	
				AI804913,	AI094960,	AI051129,	AA975822,	
			correspond to the positions of	AW367514,	AA043942,	AI337380,	AA470886,	
			nucleotide residues shown in SEQ ID	AA450210,	AA737971,	AA045559,	AL037667,	
			NO:2005, and where b is greater	AA292222,	AI914093,	AW022153,	AA620519,	
			equal to a + 14.	AA451613,	AA252687,	AAS51664,		AI953410,
				AI359851,	AA045558,	AA135778,	D58604, A	AW402976,
				AI423638,	\sim	AI189228,	AI003695,	
-				AW002772,		AI261994, D63187, AI7588	63187, AI7	58843,
				AA728996,	H02570, D	_	31974, T957	753,
				AI768841,	AW369981,		AA503361,	
				AA298895,	AI908249,		AW392006, AA962314,	
				AW392196,	AW392074,		N30487, AW392085, H52318	52318,
				AA296893,	AA303066,	AA303066, AW392190,	W35300	A031634,
				R76869, A	AA298088, T	T95752, AW3	AW391941, AI8	AI864825,
	-			Z45938, A	AA135734, N71976,	71976, AA2	AA296872, T84	T84519,
				R76870, A	AA366382, T	T81251, AA041548,	41548, C18	C18136,
				R32692, H	H02653, C16129,	129, T1082	T10828, H52227, R34	R34136,
				C17067, R	R23164, AW392168,	92168, R23	R23163, AI687114	114,
				R63893, A	AW392170, R06245, AA031753, T99872,	06245, AAO	31753, T99	1872,
				AW392082,	AW392082, AA976000, AA890237, R99970, AW238952	AA890237,	R99970, A	W238952,
				AI719088,	AI719088, AA365961, AA302997, H03271,	AA302997,	H03271, A	AA894778,
				R06300, R	R06300, R91051, D20914, W32904, AI571626	914, W3290	4, AI57162	, 6,
				AA719590,	AW386001,	AA931929,	R68979, A	AB011145,

				AR025393.	AR025401.	AR025424.	AR025397.
				AR025407,	AR025415,	AR025421,	AR025405,
				AR025404,	AR025414,	AR025423,	AR025416,
				AR025417,	AR025422,	AR025402,	AR025394,
				AR025400,	AR025413,	AR025403,	AR025418,
				AR025412,	AR025395,	AR025410,	AR025411,
		-		AR025409,	AR025419,	AR025396,	AR025408,
				AR025399,	AR025420,	AR025398,	AR025406
2006	HNHON23	889323	Preferably excluded from the	AA313697,	AA397662,	AI734131,	AA428728,
			present invention are one or more	AI734102,	AI741547,	AA428294,	AW274830,
			polynucleotides comprising a	AA428330,	AI732698,	AI742282,	AA428855,
			nucleotide sequence described by	AW452415,	AW246994,	AI337011,	AI650992,
			the general formula of a-b, where a	AA910985,	AA934713,	AW452736,	A1685505,
			is any integer between 1 to 1059 of	AW025662,	Z38485, A	AA724506, AA703833,	A703833, AA315349,
			٠,	AI653134,	AC000378,	AL080194	
			15 to 1073, where both a and b				
			correspond to the positions of				
			nucleotide residues shown in SEQ ID				
			NO:2006, and where b is greater				
			than or equal to a + 14.				
2007	HSKES11	889368	Preferably excluded from the	AI125788,	AL135619,	AI683334,	AA824310,
			present invention are one or more	AL135408,	AA037216,		AI718476,
			polynucleotides comprising a	AI829067,	W58485, A	1497128,	AW051854, N28502,
			nucleotide sequence described by	AL121373,	AI922174,	AA524333,	AW084782,
			the general formula of a-b, where a	AW402881,	AI199668,	AI199668, AI143639, AW327327,	AW327327,
				AI688325,	N42979, A	AI333116, A	
			SEQ ID NO:2007, b is an integer of	AI003784,	AI084638,	AI937411,	N29140, AW269389,
			15 to 3711, where both a and b	AA443395,	AW001384,	AI355311,	
			correspond to the positions of	AI374602,	AA424444,	AW169876,	AI335174
			nucleotide residues shown in SEQ ID	AI671042,	AW327648,	W24329,	AI050862, AI628040,
			NO:2007, and where b is greater	AA434140,	AA082441,		AI362701, AA884252,
				AI090258,	W56128, AI081404,		AA814863, W58450,
			•	AA814576,	AI907488,	AA461502,	AA223732, N95448,
				AI184687,	AI050684,	AA447362,	N21176, N27576,
				AW341550,	AA497051,	AI433749,	AA311905,
				AA505594,	AA329681,	AI278163,	AA780160,

	AA570608, AA554137, AA223721, AW271217,
	AI801216, AI097355, AA885099, W69597, AA889841,
	AI523739, AW381678, AI359091
	AA478351, AI344719, N23344, AA468529, AA235290,
	AI095678, AA608996, AI189320, AI633706,
_	AA683544, AA621614, AI857314, AI348508,
	AI214611
	AA494522, W07587, AA976842, AA478293, AA643766,
-	W46184, N35630, N33492, AA526427, N24296,
	5, N68146, H25621, AP
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-	AI869777,
	38633, T35960, AI92
	AA460576, AA383262, C17066, F06682, W16735,
	, AA876406
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	1482926, R45705, AA
	N90115,
	AI962421, AA992353, N43882, H96512, R06697,
	74107, F34857, W694
	Z19206, AA
	4, AI478732, H21536, AA318358
	61681, AW243518, CI
	, R06557, AW050504, H21535, AA
	8, AW079809, AI080026, F04276,
-	AI564126, AW070903, AW004636, H26
	AW392175, AA206629, AJ
	, AP000031, AL022336, AC00552
	AC004106, AC005015, AF001549, AC003029,

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	AC006211,		298884,	
	AC004837,			AC005229, AC007263,
	AC005	AC005011, AF165926,	926, AC005821, AL03128	AL031283,
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	U9574	U95740, AC007386,	5, Z95152, AL02	295152, AL022316, AC004990,
	AC008	AC008372, AC003070,		AL049780,
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	AC006	AC006115, AC00701	011, AL117339,	AP000694,
	AL031	AL031591, AC006101	101, AC005482,	Z84466, AC005544,
	AF001550,	1550, AC00553	ω ω	Z82206, AC005088,
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	AC006120,	6120, AC00768	586, AF053356,	AL049869,
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	AC007036	7036, AC00692	4,	AL117337,
	AP000557,		_	AE000658,
	AC005208	5208, AC005291		AL049589,
	AL022320,	2320, AC00946	464, AB016897,	AP000555,
	AP000135		, AL096791	AC002563,
	AP000563	0563, D87675,	AP000512,	U85195, AC007227,
	AL031276	1276, AC00577	779, AC005066,	U91326, Z84480,
	AL109952	9952, AC007193	193, AC005527,	AF205588,
	AC005295	5295, AC020663	563, AC005046,	AP000556,
	AC002430	2430, AC00705	055, AC004859,	AL034548,
	AL079304			AC003658,
	AC004882		AC002316	AC004841,
	AL080243	0243, Z93023,	AC007052,	Z95114, AP000210,
	AP000132	0132, AC005620,	620, AL008715,	AC006409,
	AF045555	_	-	AC003101,
	AC005	5157, AC009247	247, AP000133,	AP000211,

				AP000359.	AC004905, AC00	AC005736. AF165142.
				AC007073		757904 798036
				AC002485,		8, AP000553,
				AL049745,	AC005207, AP00045	10459, Z73358, AP000100,
				AC006013,		10
				AC009516,		AP000961, AF196779,
				AL031588,	4	AC007786, AL121754,
				AC005519,	AP000558, AP00	AP000704, AC002544,
				AC006948,	AC004491, AP00	AP000247, AC006947,
				AC006515,	_	AC002477, AC008273,
				AL049839,	284484, AC005874,	174, AF134471, AC004552,
				AL021394,	AC005839, AC00	AC005018, AC006417,
				AP000130,	AC002375, AC00	AC002126, AC005520,
				AC006992,	AC004066, AC00	AC005701, T52888, T52889,
				R06558, H	R06558, H24699, H25574, N92900, W07212	N92900, W07212,
				AA062814,	AA424971, AA93	AA424971, AA932152, AA992342, N46317,
				AA454682,	F04980, F08711,	n
				AI423663,		0262
2008	HCETP05	889467	Preferably excluded from the	AW409600,	AW370893, AW17	AW172635, H29357, H00126,
			present invention are one or more	AI688967,	H23399, H15998	H23399, H15998, AA910184, R13385,
			polynucleotides comprising a	AI635135,	AA811899, AA76	AA811899, AA768537, AA827197,
-			nucleotide sequence described by	AA152215,	T33955, AA324892,	92, H51900, AW015309,
			the general formula of a-b, where a	Z45802, A	AW138603, AW439297,	97, AA281159, T31539,
			is any integer between 1 to 454 of	AI989451,	AA311444, T33897,	
			SEQ ID NO:2008, b is an integer of	AL096745,	AL133562, AB02	ហ
			15 to 468, where both a and b			
			correspond to the positions of			
			nucleotide residues shown in SEQ ID			
			NO:2008, and where b is greater	·		
			than or equal to a + 14.			
5000	HDHEA53	889494	Ω	AW162106,	AI192344, AI564803	4803, AI816163,
			present invention are one or more	AW157769,	W30860, AW157220,	20, AI686640, AI379866,
			polynucleotides comprising a	AI917170,	AA548108, AI58	AIS81151, AA190572,
			nucleotide sequence described by	AA479158,		AI827282, AI400087,
			the general formula of a-b, where a	AI271370,	AA609367, AA23	AA236262, AI910788,
			is any integer between 1 to 825 of	AI148957,	AA758679, AI392976	2976, AA608963,

			SEO ID NO:2009, b is an integer of	AA464601, AI634775, W07097, AI332514, AA253390,
			Ø	AA490370, H80788, AI024529, H21486, AI338291,
			correspond to the positions of	N72328, AA193686, AA253494, AI240331, H20219,
			nucleotide residues shown in SEQ ID	AA064633, AA664481, AA548109, R10906, R61486,
			NO:2009, and where b is greater	AI970230, AI652083, AI654228, H75492, AA247266,
			than or equal to a + 14.	N52829, AW139159, AA748177, R64411, H17572,
				AF065389, AF053455, AF121344
2010	HCHAC08	889700	Preferably excluded from the	4
			present invention are one or more	AI743223, AI804911,
			polynucleotides comprising a	AI034362, AA468381, AI168829, AA468421,
			nucleotide sequence described by	AA860298, AA578670, AI027557, AI365637,
			the general formula of a-b, where a	AA618558, AI307591, AI033866, AA052982,
			is any integer between 1 to 799 of	AA937189, AI034209, W05444, AA612975, AA053475,
			SEQ ID NO:2010, b is an integer of	AA468294, AI972035, AA612979, AW004657, N58184,
			15 to 813, where both a and b	AA782754, AI186935, T53519, AW016322, R27278,
			correspond to the positions of	AA988007, AA579074, AA860739, AA612976,
			nucleotide residues shown in SEQ ID	AW406518, AI422596, F25986, AA774165, N56542,
			NO:2010, and where b is greater	AA864684, AA922471, AA468220, AI350544,
			than or equal to a + 14.	AI950616, AI142741, AA706997, C21238, T53520,
				AA095378, AI673154, AI905956, AI660174, T24673,
				F36466, AI341288
2011	HACBT96	889782	Preferably excluded from the	AI338644, AI745184, AI890849, AW079838,
			present invention are one or more	AW149663, AI634926, AI889135, AW026717,
		_	polynucleotides comprising a	AW270045, AI857571, AI052517, AI004249,
			nucleotide sequence described by	AI279282, AW089862, AI499010, AA581431,
		_	the general formula of a-b, where a	AA669174, AW129569, AW438690, AA830692,
				AA419072, AI624275, AA434407, AI597766,
				AI184077, AA565719, AA758787, AI183979,
			15 to 994, where both a and b	AW021522, AI862132, AA705896, AI090447,
		_		AA828220, AI190867, AA435546, AA568841,
			nucleotide residues shown in SEQ ID	AI092061, AI146792,
			NO:2011, and where b is greater	AA012947, AA700657, AI160133, W90656, AA618520,
			than or equal to a + 14.	AA805610, AL043849, AA902677, AI276955,
				AI366145, AA394012, N74351, AA076429, N92748,
				N74405, AA830815, AA788867, W86234, AI131041,
				AI636459, AL043850, AI309739, AI346161,

			C 1 1 0 4 4
			0, AMICUACO, AIGOROS, AMOUTOCO,
-			y · w
			O NICKES, MACHOUS, MISTOSOL, MWCKOLS
			8, ALU45358, AA669031, H81440, F25
			, H41079, AA433970, T03708, F37029,
			, AA923050, W67263,
			AA604066, AI186384, T55826,
			AI568300, R07512, AA862409, AI350206, R44871,
	_		H46287, AA857126, AI
			T74684, H75881, D25565, AA419133, AW188884,
			, AA3058
		-	, Н66896
			, T95777, AW384420
			R12501, C21226, AI547271, R07565, AA326036,
			AI610783, T95776,
	-		ന
			E72070, AA
			AA345444, T74921, AA404700, R08428, AA934685,
-			, C04482,
			U46347,
			, K03001
			M20456, M2
			M20455, AF164120, AR060234, AB026436, U02317,
\dashv			AR066494, AR069079, AR054110, A81671
2012 HTLEN01	889954	Preferably excluded from the	T08846, AA
_		present invention are one or more	AW072169,
		polynucleotides comprising a	
			AA535028, AI139078, AA077934, AI361426,
		egeneral formula of a-b, where	AI359977, AW009454, AB033050, AB015330
		teger between 1 to 1756	
		SEQ ID NO:2012, b is an integer of	

and b ons of of in SEQ ID greater	n the more ng a ribed by 1-b, where a to 693 of integer of and b ons of n in SEQ ID greater	AA878377, AW264482, AA528458, AI084502, le or more AI086537, AA280756, AI524467, AA215387, al909056, D20028, AI432571, T80449, C16437, al474660, AA306817, AA636097, AA214516, R82222, arb, where a AA995304, R39369, AA318653, R62525, AL045794, to 2426 of AL03924, AA969711, D51250, T24119, T24112, integer of AL039629, AL039648, AL039726, and b AL039629, AL039625, AL039648, AL0395837, and b AL039074, AL039678, AL039108, AL039538, an in SEQ ID AL039509, D80219, AL039539, AL039556, greater AL039109, AL04992, T80169, AL04530, AL039423, AL044407, AL038821, AL039410, AL0433422, AL04445, AL038025, AL039410, AL043442, AL043445, AL038025, AL03941, R24660, D51423, D80134, D59619, AL043441, R24660, D51423, D80134, D59619, AL0469, D80196, C14227,
IP 0 : >	preferably excluded from present invention are on polynucleotides comprisinucleotide sequence describe any integer between 1 SEQ ID NO:2013, b is an 15 to 707, where both a correspond to the positinucleotide residues show NO:2013, and where b is than or equal to a + 14.	Preferably excluded from present invention are or polynucleotides comprision nucleotide sequence describes any integer between 1 SEQ ID NO:2014, b is an 15 to 2440, where both a correspond to the position nucleotide residues show NO:2014, and where b is than or equal to a + 14.
	889962	88 999 4
	HCROA43	HSLJW05
	2013	2014

	AL039085, AL036196, D59927, AL037639, D80949,
	D80366, AL037615, AI535783, AW451070, D80168,
	., AL036767, AW452756
	T11051, D50995, C75259, C14014, AL037526,
	D80045, R47228, AL037104, AL036679, AI910647,
	AL036924, AL037601, D59889,
	AL036733, C15076, AL036158
	', T23659, AL036418, D80038,
	AL037054, AL036765, D5828:
	D80188, AL037177, D59467, C1
	10378, AL037081, AI
-	98, ALO37047, F13647, ALO37643
	, T48598, AL036964, D50979,
	D80522, AA514190, D80212, AL036132, AL036167,
	AL037600
	AA285331, D59502, Z21582, AL037679, D80164,
	D80166, AW
	AL036152,
	,042628, AI
	2, AI287326, D80024,
	L039086, AL
	AI591316, AI499285,
•	, AL043326, AI955906,
	, AA225339, AI763414,
	6, AW150578, AI538085,
	5, AI857296, AL045163,
	5, AI815855, AI687127, AI86657
	5, AL119791,
	8, AI174394, AI612885,
	, AI252023,
	8, AI620284,
	, AW082113, AL042745,
	AW089572, AI433976,
	L045500, AI866770, AL022401, A8539
	A85477, AR025207, X68127, A86792, A44171,

	A67220, I18371, AR062871, AR037157, AR017907,
	A84772, A84773, U87250, AR062872, A84776,
	A84775, AR062873, A84774, AR067731, A20702,
	A58522, A91750, AR067732, A43189, A43188,
	A38214, A98767, A95117, I56772, I95540,
	A93963, A93964, A51047, A63064, A18053, A49700,
	AR018923, AR031375, A48774, I63120, A63072,
	A75888,
	A60111, A23633, AR015960, A23998, AR000007,
	AR020969, I03343, AR054109, I06859, AR022240,
	A81878, A64081, A58524, E12615, A24783,
	AR035193, A24782, A58523, A92133, E14304,
	E16678, A82653, E16636, A93016, AB012117,
	Z96142, I26929,
	A58525, A02712, AR038762,
	V00745, I49890, AR000006, I19516, I44516,
	3, E13740, AR008430, Y11923, AR03
	A11245, A02710,
-	A13392, I19517, A76773, A22413,
	Y17188, A35537, A35536, A97211,
	A02135, A04663, D28584, A51384, I
	I08051, A70040,
	AR035974, AR035977,
	4, AR038286, I66495, I664
	I66496, I66494, I66486, I66487,
	I03664, E00523, I92483,
	14548, A10361, I00077, I19525, AF
	65, AF019720, S70644, I07429, E0603
	A18722, D26022, X13220, AF156304, A91754,

				AR027069, A20701,	A20701, A04710, A52326, AF096810,
					X58217, A62298,
				A60957, I	I84554, A62300, I84553, S65373, A60968
					AF096793, A60985, A60990, A60987,
				I69350, A	A84916, Z82022, D44443, AB007195,
					A80951, A10363, AR018138, AF156300,
				X73003, A	AR028564
				vo	E04616,
				A08457, A	AJ132110, S69292,
				A13038, A2	
				I07888	
2015	HTPGK74	89068	Preferably excluded from the	AI149400,	AA846733, AI085373, AI246729,
			present invention are one or more	AI608911,	AI923892, AI798918, AW303427,
			polynucleotides comprising a	AI708285,	AW080676, AI684195, AI587306,
			nucleotide sequence described by	AW189579,	AI354582, AW044409, AI922230,
			the general formula of a-b, where a	AI628502,	AI888388, AI758885, AI619483, W52043
			is any integer between 1 to 3288 of	AW057673,	AL037160, AI921372, AW304335,
			SEQ ID NO:2015, b is an integer of	AI624382,	AI819541, AW276527, AI554494,
			15 to 3302, where both a and b	AI809216,	AI923339, AI381549, AI015540,
			correspond to the positions of	AI473800,	AW104317, AI910909, AI471516,
			nucleotide residues shown in SEQ ID	AI624577,	AI141307, AA075786, AI807993,
			NO:2015, and where b is greater	AI566219,	AI589224, AL048943, AI620365,
			than or equal to a + 14.	AW391429,	W37101, AI346763, AW086487, AI457487
				AA664049,	AW440483, AI476665, W63597, AI254213
				AA934370,	
				AI469784,	, AI911889,
				AW439282,	AI038691, AA884808, AI798465,
				AA905923,	AA075733, AW372900, AW385528,
				AW385522,	AL119459, AI352172, AA587707,
	•			AA947323,	AA732669, AA617672, AI282902,
				AA946660,	AI342280, AW372893, AI589501,
				AI185284,	N90643, AW363861, AI051922, AI963833
				AA583025,	AW372901, AI221733, AI922401,
				AA912170,	
				AI952654,	AI087392, AI565591, AA931720,
				AI026053,	AA838395, AI361077, W28135, AI025960

	AI458833, W28630, AI539757, AA532831, AI272036,	, AI254219, AI025959, N23305,	, AA999740, AI369031,	AI084385, AA846191, AA721128, AA463497,	AI278750, AA767535, AA91324	AA649145, AA115785, AA904917, AI953598,	AW004980, AI433221, AI167837, AA878628,	AA725431, AI311013, AI358579, R19761, W39658,	AA164801, W19891, W73947, AI955376, AA872844,	Ξ.	AA164725, AI188067, N34986, AA428266, AW166792,			AI682818,		N95218, AW272360, H55826, R83098, AA605309,	2, AA648856, W38889, AA70	AA115311,	H55818, R86911, AA305294, R94357, AI914666,	AW373543, W21046, AI926759, C06443, H47177,	. M31393, T10966, H74049, H61903, N32148, R35482,	AI565915, R86899, H71088, AA021144, H13058,	AW009569, AI630631, AA336531, AA568673, D58749,	,	AA970031, AI914563	21095, AI557184,			R66061, T77140,	AI290392, R33973, R64607, X59408, X59405,		A18585, AR063631, X59409, X59410, AR063630,		AR066587, D84105, S51940, D85750, E05680,	1000011 0000011 VECTOR 0101VA 1101VA VIENOR
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				AF067728, Y14314, E0 AL137527, A45787, AF AF118070, AF162270, T85655, R0 R22036, R2 R40303, R4 R70795, R7 H08122, H1 R83039, R8 H96583, H9 W31430, W3 AA035520, AA164800, AA689363,	Y11587, 2221, AF AJ006417 061943, 272491, TT55602, 15955, R0 (2364, R2 (2364, R2 (2364, R2 (3150, H1) (6755, H5 (6704, N4 (1986, N5 AA079896 AA079896 AA079896	AL133081, A93. 118064, X87583. 1 AR013797, IC AR013797, AL0. 282022, AL1177 10661, R13220, 17810, R3367, 17810, R64594, 13260, H04294, 13260, H04294, 13260, H0598, 1238, AA02117, 1238, AA02117,	AL133081, A93350, AF02612 118064, X87582, AF079763, AR013797, I03321, AL080 AR013797, AL049283, AL050 Z82022, AL117585, AF12594 T55684, T83368, T83513, 6061, R13220, R15276, R21 7810, R33367, R33366, R84 5180, R64594, R66060, R69 10960, H04294, H04562, H08 3266, H26621, H26673, H47 5819, H55827, H71875, H73 9329, N71908, N93708, W24 1238, AA021171, AA035032, AA079897, AA165060, AA225869, AA632555, C01059, AA091046, AA09133	026124, 9763, ALO80159, ALO50024, 125949, 13, R21850, R34281, R69585, H08121, H73487, H73487, H73487,
2016	HHGAB64	869068	preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 365 of SEQ ID NO:2016, b is an integer of 15 to 379, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:2016, and where b is greater than or equal to a + 14.	AA127776, AA477584, AL031295,		φ _φ	0	T67843, F022382,
2017	HOSOR86	890753	preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 2042 of	AI341460, AI909124, AI801242, AA974210, AA907244,	AW173384, P AW118938, P AW438695, P AW130020, P AI146982, P	AWO55235, AI689438, AI123971, AA489046, AI093766, AA983814,	W39204, Al AI419443, AA707755, AW298736, AA284319, AI955386,	AI909118, , N59864, , N59886,

				AI859864, AI498376, W01363,
			\mid 15 to 2056, where both a and b	AA824487, T86598, AA994605, AW044013, AA489144,
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	T77559, AA736753, T77523,
			NO:2017, and where b is greater	H44608, AI955411, N90263, H94626, AL119283,
			than or equal to a + 14.	AL119309, AI909117, N77027, N79005, AW105078,
				N62828, AI334730, AI701272, T07505, AW376940,
		,		AI909110,
				06840,
2018	HE9RV77	890763	Preferably excluded from the	AW241738, AIS54315, AW293947, AI763258,
			present invention are one or more	AI721194, AW043707, W67989, AW269975, AW025268,
			polynucleotides comprising a	AI683778, AW183594, AW242994, AW015541,
			nucleotide sequence described by	AA428411, AI312039, AA905967, AW274692,
			the general formula of a-b, where a	AI743918, AI632220, AA515764, AI018660,
			is any integer between 1 to 1877 of	AA936423, N40612, AI913282, N36286, N42415,
			SEQ ID NO:2018, b is an integer of	AA155820, AA155924, AA071299, W68001, AI799025,
			15 to 1891, where both a and b	
			correspond to the positions of	_`
	•		nucleotide residues shown in SEQ ID	AW168113, H64050, AI261230, AI347397, AA536165,
			NO:2018, and where b is greater	AI569491, AW172624, AA781882, AI583725,
			than or equal to a + 14.	AA149663, AA683414, AI539802, AI583700,
				AI445057, AI816810, AA176623, AI340128,
				AA375927, AA628568, AA434428, AA164797, R80702,
				AA445933, AI690654, H10573, AA179678, H15588,
				AI313391, AI538861, AI687194, AA167315, N25332,
				AW150559, W58766, H17389, AA164796, H82362,
				:204281, AW301352, AW302888,
				AA218953,
				3, AA151878
				AI282274, TE
				AA173300,
				R11810, Z39244, T71190, R17172, R17252, N24523,
				\sim
				9, R14564, AA102051, T81858, AA
				D31565, U46380, AI277142, AA628822, F06639,

				AW004021, AA860192,	AI500444, H61486, H87106, AI254025,		AI962340, AI675481, F04003, AA166985,
				AA321073,	AISS7191, A		AW072197,
				A19221/1, AW168889,			
				AA090327,	AI218075, AA383806, AA220919	AA383806,	AA220919,
				AA102050,	H10369, AF	133426, AF	H10369, AF133426, AF053453, AF043906,
				U84895, Al	U84895, AL035608, AF053454, AT633192		D16949, AI336283,
2019	HPRAJ70	890776	Preferably excluded from the	AI805082,	AI432462, A	AW263421,	AA135870,
			present invention are one or more	AA137165,	AA298464, 1	AA298471,	AA298475,
			polynucleotides comprising a	AA298489,	AI362575, A	AA031604,	AA313094,
			nucleotide sequence described by	AA031360,	AR009514, 1	AF079864	
			SEQ ID NO:2019, b is an integer of				
			15 to 3557, where both a and b				
			correspond to the positions of	•			
			nucleotide residues shown in SEQ ID				
			NO:2019, and where b is greater				
			than or equal to a + 14.				
2020	HBODK52	890801	Preferably excluded from the	AI554661,	_	AA314190,	AL120376,
			present invention are one or more	AI334374,	AI274093, 1	AI080270,	AA883816,
			polynucleotides comprising a	AA879435,		AI222322,	AI432982,
			nucleotide sequence described by	AA541454,		AA749031,	AA307355,
				AA993688,		F24838, AI	F24838, AI147394, AI864022,
			is any integer between 1 to 1585 of	AA298719,	7	~	~
			SEQ ID NO:2020, b is an integer of				
			15 to 1599, where both a and b	Z44194, A	AW139211, AL	AL008582, AE	AB035207, D64109,
			correspond to the positions of	AL022393			
			nucleotide residues shown in SEQ ID				
			NO:2020, and where b is greater				
			than or equal to a + 14.				
2021	HARNK52	890820	Preferably excluded from the	AW372332,		AW372303,	AW392509,
			present invention are one or more	AW392497,		AW372464,	AW392505,
			polynucleotides comprising a	AW004891,	AA101225,	AW392512,	AA102670,

			nucleotide sequence described by	AA120821.	U54597. A	AW182872 AT446810	ŀ	8798878
			the general formula of a-b, where a	AA294978,	AW392492,	AW392492, AA298897, U54599,		AI903382,
			is any integer between 1 to 2579 of	AA991253,	U95367, I	U95367, I59650, U95368, AF009702	0	702,
			SEQ ID NO:2021, b is an integer of	AF009697,	AF009701,	AF009700,	AF009699,	
			15 to 2593, where both a and b	AF009695,	AF009693,	AF009694,	AF009698,	AF009696
						•		
			ಌ					
								-
			than or equal to a + 14.					-
2022	HTLHU22	890863	Preferably excluded from the	AW248608,	AI654134,	AW249047,	AW027462,	
			present invention are one or more	AI688329,	AW136847,	AA995019,	AI867957,	283847,
			polynucleotides comprising a	Z82206				
			nucleotide sequence described by					
			the general formula of a-b, where a					-
		-	is any integer between 1 to 1674 of					
			SEQ ID NO:2022, b is an integer of					
			15 to 1688, where both a and b					
_			correspond to the positions of					
			nucleotide residues shown in SEQ ID					
			NO:2022, and where b is greater					
			than or equal to a + 14.					
2023	HWMBB2	890945	\overline{C}	AL042015,	AI760156,	AI041208,	AI675831,	
	6		present invention are one or more	AA772287,	AI761091,	AA127766,	AI189553,	
			polynucleotides comprising a	AI024414,	AI680106,	AA678819,	AI338208,	
			nucleotide sequence described by	AI276652,	AA069849,	AI457552,	AI005201,	
			al formula of a-b, where	AA678586,	AA918062,	AA411763,	AA037163,	
			teger between 1 to 2529	AA069802,	H30857, A		AA216712, A	AI266630,
			SEQ ID NO:2023, b is an integer of	N23150, AI082636,			AA385301, AJ	AA411843,
			15 to 2543, where both a and b				AA347097, T	T28624,
			correspond to the positions of	N32729, AA	AA146702, A	AA343535, AA	AA375419, AV	AW316863,
			nucleotide residues shown in SEQ ID	N32133, AA	AA385302, A			AA375420,
			NO:2023, and where b is greater	AI867611,	AW206128,	AI867611, AW206128, AI630096, N95166,		AA331777,
			than or equal to a + 14.	Z24775, AA	AA331778, F	F04253, AA318183,	05	300,
				F04964, AA	1343617, A	AA343617, AA194918, R41937,	11937, AA3	AA347119,
				AI524404,	AA362621,	AI524404, AA362621, AA402478, F00058	F00058, A	, AW366370,
				C21140, R1	.0662, ALO	C21140, R10662, AL079560, AA994433,	94433, AA2:	AA218592,

				AA888498, AA371095, AI783880, AA293830,
				U07343, UC
				U40971, U40978, U17857, U40975, U17854, U17849,
				U17851, U40972, U17839,
				U17847, U40977, U17852, U40968, U17841, U40973,
				U17844, U40976, U40962, U17855, U40965, U17846,
				U40961, U40967,
_				U17842, U40963, U40964, U17843, U17853, U40974,
				S77856, U17845, U40966
2024	HWLND63	891125	Preferably excluded from the	643, S750
			present invention are one or more	E02516, M37721, E03981, AF010472, E03204,
			polynucleotides comprising a	E03203, E03201, D29625,
			nucleotide sequence described by	M18683, E03205, I09286, U79523, M82845, E03428,
			the general formula of a-b, where a	AR036184, X59689, M25719, M25732, X59687,
			is any integer between 1 to 490 of	X59688, E02517, X59685, X59686, T47438, T49517,
			SEQ ID NO:2024, b is an integer of	T40337, T41197, T94036, R31007, R52165, R54705,
			15 to 504, where both a and b	
			d to the positi	H09249, H13692, H13744, H14286, H20221, H24797,
			nucleotide residues shown in SEQ ID	H25936, H25967, H27194, H27195, H27531, H28158,
			NO:2024, and where b is greater	H30301, H42178, H39094, H43206, H43253, H43704,
			equal to a + 14.	H43788, H43842, H44053, H44129, H46393, H47935,
				R87926, R89640,
				H84491, H93855, H95554, H96000, H96001, N29623,
				W20057, W56622, W56652, W73707, AA001437,
				AA001129, N91455, N91545, AA010455, AA012908,
				AA017259, AA017548, AA019579, AA021397,
				AA021267, AA031311, AA031448, AA054148,
				AA055244, AA055263, AA057094, AA079530,
				AA079578, AA086369, AA086477, AA086052,
				AA088887, AA088908, AA101239, AA112044,
				AA112875, AA113195, AA113785, AA121382,
				AA134323, AA134324, AA134404, AA134405,
		· · ·		AA159956, AA159957, AA169782, AA179024,
				AA179789, AA190506, AA190992, AA191267,
				AA191540, AA193244, AA194300, AA194320,
				AA194750, AA194569, AA195818, AA196755,

10, 10, 35, 32,	8, F15660, 6, F16798, 2, F17509, 063,	8, 3, 5, 0, 9, 718024,	8, F18564, 5, N84794, 3, C02843, 5, C03831, 964,	8, 0, 5, 5, 5, 6, 9,
AA235645, AA24330 , AA250964, AA25094 , AA459644, AA46416 , AA427566, AA48033 , AA533290, AA55522	AA558632, AA5639 089, F16376, F165 260, F17364, F174 566, F17588, AA58 AA583809, AA5839 AA594803, AA6042	AA657777, AA65798 AA806213, AA82754 AA857063, AA86553 AA876266, AA91741 AA962483, AA96886 AA975454, AA97639	117, F18383, F1841 190, F19528, F1971 115, W73754, N8922 145, C03180, C0332 141, C05199, AA018 AA642243, AA09594	15, AA213946, AA28498 129, AA291918, AA29200 62, AA293262, AA40190 F20840, F20860, F2151 118, AA456784, AA45451 02, AA477416, AA47774 15, AA481936, AA48146 76, AA628543, AA66637
AA197162, AA223624 AA250844, AA250903 AA459406, AA459418 AA464690, AA464779 AA483686, AA508610 AA515092, AA515572	H H H	, 10 H 4 L 9 E	11100	AA554007, AA211715, AA284536, AA290829, AA293474, AA293062, F20441, F20482, F20 AA411329, AA410818, AA459631, AA47102, AA477852, AA480115, AA45654, F21911, F

				F22724, AA719223, AA724815, AA725731, AA758587,
				5, AA812572, AA845555,
				, AA852552,
				AA993537, AI025737, AI038538,
				97, D256
_				F00386, F01041, F01120, F01124, F01135, F00308,
				AA77
				AI347597, AI361314, AI361315, AI361322,
		_		AI401660, AI423575, AI423596, AI128394,
				AI224046, AI144391, AI149311, AI625219,
				AI625399, AI192566, AI214910, AI658645,
				AI538037, AI342442, AI633128
2025 F	HCROQ71	891264	Preferably excluded from the	Z99396, AW392670, AW38
	,		present invention are one or more	
			polynucleotides comprising a	AL119319, AL119396, AL119457, AL119324,
	•		nucleotide sequence described by	AL119341, AL11939
			the general formula of a-b, where a	AL119355, AL119496, AL036418, AL038837,
			is any integer between 1 to 766 of	AL119335, U46350, AL119522, U46349, U46351,
			SEQ ID NO:2025, b is an integer of	AL037051, AL036725, AA631969, AL042970,
			correspond to the positions of	H
_			Ð	AL037205, AL039074, AL119439, AL036924,
			NO:2025, and where b is greater	AL042544, AL038509, AL042975, AL119488, U46345,
			than or equal to a + 14.	AL134538, AL042984, AL042551, AL134527,
	_			AL043029, AL042542, AL042450, AL037094,
				AL037526, AL037085, AL036196, AL037082,
				AL037639, AL037077,
				AL043003, AL036767, AL036190, AL036268,
				AL038520, AL038851, AL119464, AL038447,
			_	AL036774, AL036998, AL036733, AL037178,
				AL036238, AL037615, AL037027, AL036719,
				σ
				AR
				AR064707, AR069079, AR054110, AB026436
2026	HBINP81	891305	Preferably excluded from the	AI206965, AI955864, AI978772, AI952843,

polyn nucle the g is an SEQ I SEQ I 15 to corre nucle No:20 than SEQ I SEQ I than oucle No:20 than	891896	HDLAG89 891896	ent invention are one or more nucleotides comprising a eotide sequence described by general formula of a-b, where a ny integer between 1 to 2507 of 1D NO:2026, b is an integer of o 2521, where both a and b espond to the positions of eotide residues shown in SEQ ID 026, and where b is greater or equal to a + 14.	erablent in nucle entrol entro
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AI538885, AI431230, AI889147, AL041862,	AL042515, AL045500, AL046356, AI433976,	AI872300, AL042551, AW172723, AI440263,	AL039390, AI371251, AI866510, AI436429,	AI371228, AL040207, AI890907, AI860003,	AI610557, AI866465, AI887499, AI431321,	AI690946, AL045328, AI866469, AI521594,	AI828574, AL048427, AL042538, AI537515,	AL043089, AI275175, AL042745, AL043091,	, AI648567,	AI499463, AI582912, AI610362, AI538850,	AI887775, AI623736, AI590043, AL045620,	A1440239, A1492519, A1539800, A1923046,	AI434242, AI500714, AI537273, AI355779,	AI885949, AI581033, AI491710, AI436456,	A1469775, A1963846, A1567940, A1817244,	AI612913,	AI671642, AI285826, AI539707, AI863014,	_	AL046681, AI521571, AI432677, AI610357,	AL042377, AI434223, AI366900, AI610429,	AI539632, AI889148, AI539847, AL042939,	AIS67935, AI805762, Z98465, AIS61170, AI702065,	AI354998, AL047422, AL045891, AI344785,	, AI866820,	, AW172745,	AI433968, AI567953,	AW403717, AL048656, AI866461, AL047092,	AA420758, AI521465, AL043321, AL039276,	AI371265, AI049851, AI274759, AI866457,	AI285419, AI927233, AI567993, AI431315,	AI654276, AI628850, AW118237, AW191003,	AI828583, AI539863, AW162194, AI364788,	AL045163, AL110306, AL048323, AI521596,	9108, AI554827, AW197139, Y17793, A93	AR066494, A58524, A58523, AL137429, U77594,

	AF090901, AL133049, AL050116, AF091512, E05822,
	Y11587, AL122049, A08916, AL137539, L10353,
	I48978, A08910, A08909, AF100931, AC004883,
	, AF182215, E07108, AC004227,
	AF113694, AJ000937,
	AL122110, I89931, AF118090, AL080124, AF111112,
	X65873, AL050108, X89102, AB03
	L137271, AL137521,
	AF118070, AL122050, I48979, AF090896, AF100781,
	Z72491, AL137538, Z37987, X83508, A65341,
	6, AF158248, AF177401, S687
	E00717, E00778, I26207, AF097996, AL133080,
	AL137459, AC006840, AF102578, E01573, E02319,
	E07361, A57389, AL049430, A90832, A93350,
	Y16645, AC
	AL133557, AL137529, Y07905, AL133560, AL137463,
	AF119337,
	AL049283, A08908, I33392, AI
-	AL117585,
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	m
	U78525, AL117435, AL12
	AF113019, AL137283,
	AL110196, AF087943,
	AL137658,
	A18777,
	AL049382, AF125949, AL096744, AR034821, A03736,
	AL122121, U72620, AF081195, AF111849, Y10655,
	X70685, U42766, U58996, Z82022, AC006313,
	AJ242859, AF183393, AL050149, AL110225,

				AL122118. AF106862. U88966. AF026816. A18788.
				, AF067790,
				AL110197, AL137648, AL117460, AF026124,
				AF090900, AF125948, AF039138
				AF039137, AL133014, AL117394, A12297, AL133031,
				AF079765, X63574, X96540, AL110280, X98834,
				AR020905, AL137556, I09360, AL133093, AF067728,
				AF113699, AL137560, I42402, L31396, AL133568,
				AL137488, AL050138, AL050393, L31397, U91329,
				AJ012755, AC006039, I89934, X52034, AF126247,
				X84990, AL080127, AL133075, AJ003118, AL080137,
			-	AL137527, AB029065, AF113691, AF061943,
				AL137476, S75997, X94372, AR013797
2028	HE8FL95	892113	Preferably excluded from the	AA195218, AA397579, AA399552, AA621184,
			present invention are one or more	AI692940, AA205886, AI702167, AI365354, AF090947
			polynucleotides comprising a	
			nucleotide sequence described by	
			the general formula of a-b, where a	
			is any integer between 1 to 1769 of	
			SEQ ID NO:2028, b is an integer of	
			15 to 1783, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:2028, and where b is greater	
			than or equal to a + 14.	
2029	HHFGIS9	892177	Preferably excluded from the	AW207619, AA534290, AW340566, AW139543,
			present invention are one or more	AA947281, AA776464, AI697902, AA037301,
			polynucleotides comprising a	AA205320, AA443876, AI206904, AA400700, T75075,
			nucleotide sequence described by	AI363369, W46782, H24404, AI032106, AI880884,
			the general formula of a-b, where a	N59387, AI138757, AA307337, AA554317, AI359282,
			is any integer between 1 to 4317 of	N28440, AW007847, AA662978, AI129939, AA476728,
			SEQ ID NO:2029, b is an integer of	AI017751, AI431939, AI675507, AA953932,
			15 to 4331, where both a and b	AI625227, AI991609, H23505, H18538, N77075,
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	AA282393, F02661, AI635585, AA024899, R80487,
			NO:2029, and where b is greater	T77003, AI040191, AI363266, H14797, H68321,

	than or equal to a + 14.	AA024900, R83449, AI249693, Z42220, AI560382,
		AI301618, Z43207, R80381, AA20675
		F06352, W93287, R40397, T87366, AI767771,
		AI094857, F02642, AA970085, AI942231, F06371,
		F12724
		10631,
		50637,
		AA331899, AA307511, AW363028, AA296346,
		AA400655, F07384, T98853, R13009, AW169922,
		AA218742, AI827798
		.3181, AI
		AI023953, AW316878, D80045, F11062, AA581647,
		AI382497, D59502,
		C14429, D58283, D81030, D80195, D80043, D80227,
		C14331, D80188, D80038, D51423, D59619, D80210,
		D51799, D80391, D80240, D80253, T03269, D80166,
		3, D80196, D80269,
		, D59275, D57483,
		Z21582, D80378, D80366,
		D50995,
		275259, C1
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		AI557751, D80268, AA305578,
		D51079, D51022, AW179328, AW177440, AW178775,
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_		C14407, C14227, AW
		A
		AW369651, AW178762, AW177501, AW177511, D51097,
		, D80133, AW360811, C1
		, AW375406,
		, AW377671, AW360834,
		, AW378534, AW179332
		23, AW178905, T11417, D80302,
		D80132, AW177505, AW352171, D80439, AW377676,

AW178906, AW352170, AW177731, AW178907, AW179019, AW179024, AW179220, AC005534, AC007075, A62298, A62300, A84916, Y17188, AJ132110, U87250, A82595, AR018138, A78862, D26022, A67220, X67155, D89785, D34614, A25909 X82626, D88547, AR008278, AF058696, I19525, AR025207, AB028859, X68127, A94995, I82448, A44171, AR016808, AB012117, Y12724, A85396, AR3190, AR060385, AB002449, A30438, AR008443, A73190, AR060385, AB002449, A30438, AR008443, A71187, AR038669, AR008277, AR008281, AR066488 AR060138, A45456, A26615, AR052274, I18367, Y09669, A43192, AR066487, I14842, AR054175, D88507, AR066490, D50010, AB023656, U79457, AB033111, U46128, AR064240, A63261, AR016691, AR016690, AR008408, AR062872, A70867, D13509, I79511, A64136, A68321, AR060133, U87247, AF123263, Z32749, AR032065, X93535, AR008382	ALO36113, AA433879, ALO45190, AA057554, AW239170, AW403966, AW239410, AW402407, AW375966, AW373031, N20475, AW067770, AW179034, ad by AA410697, AA074710, AI752785, AW068103, H10878, where a AI869324, AW067904, AI751896, R87863, AA603295, 1220 of R56461, W86435, AW068684, R88501, N25503, ager of A1909381, M78217, AA852669, R87623, R87854, ab AA018954, AA410887, AA057207, AA362685, R84663, of A8018954, AA410887, AA018988, T87278, H01627, ater R85220, AW388463, R85353, AA35125, AW076608, AW062971, AA852517, H72226, AW393764, W02259, AW062971, AA852517, H72226, AW393764, W02259, AN062971, AA852517, H72226, AW393764, W023337, K699954, H27662, R24815, T99262, H52007, M11233, X05344, X52886, X53337, M63138, M63137, M63135,
	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where is any integer between 1 to 1220 SEQ ID NO:2030, b is an integer of 15 to 1234, where both a and b correspond to the positions of nucleotide residues shown in SEQ NO:2030, and where b is greater than or equal to a + 14.
	892291
	HOFMT75
	2030

HWLEQ37	892367	Preferably excluded from the	AI884627,	AW130437,	AI668781,	AL043335,
		present invention are one or more	AI907430,	AI613418,	AW150279,	AI205012,
		polynucleotides comprising a	AI038777,	AA709407,	AI690430,	T35506, AI022430
-		nucleotide sequence described by	AI023459,	AI476713,	AI671575,	D29621, T34271,
		the general formula of a-b, where a	AW419081,	AI261913,	N32030, AA377446,	A377446, AI912514
		is any integer between 1 to 1075 of	AA034072,	AI053445,	AI828656,	AA533408,
		SEQ ID NO:2031, b is an integer of	AI038724,	AL042113,	AI370475,	AA569743,
		15 to 1089, where both a and b	AI623899,	AL135698,	AI283090,	AW272763,
		correspond to the positions of	AI868164,	AA633266,	F17700, H	F17700, H57826, A1633185,
-		O	AL045709,	AA713674,	AA360944,	AA716755,
		NO:2031, and where b is greater	AW088125,	AA297968,	AA659324,	AI252506,
		than or equal to a + 14.	AB020865,	AC005940,	AP000694,	299755, AP000557,
			AL035587,	AC004701,	AC006965,	AC005768,
			AC006211,	Z98950, AC	AC005152, Z8	85996, AF051976,
			AL109963,	AC007934,	AC005339,	AF053356,
•			AC006116,	AL031281,	AC005755,	Z69917, AC005599
			AC003101,	AL050308,	AL096791,	AC005288,
			AL132992,	AC004472,	AL022326,	AC004386,
			AL021368,	AP000073,	AP000512,	AC004148,
						AC002527, AC005821
			Z99714, A	AC006387, AC	AC002375, A	AC006547, AC005041
			AC006285,	AC002347,	AC002045,	AC002418,
			AC003010,	AC005520,	AP000248,	AC005192,
-			AC005225,	AP000346,	AF001549,	AC006480, Z93017,
			AC005899,	AL096801,	AL096817,	AC003982,
			AL121652,	AC005189,	AC005968,	AC005212,
			AL035683,	AL049829,	AF196779,	AL022320, L44140,
-			AC005971,	AL022721,	AC005701,	Z85987, AL133448
			AL109865,	AL031055,	AC004236,	AC000097,
			AC006026,	AC004682,	AC004894,	AC005015, 299716
			AC001228,	AL133245,	AL035400,	AB023049,
			AL031286,	AP000279,	AC004797,	AC007226,
			AL139054,	AP000260,	AL034429,	AF111168,
			AB023050,			C006046, AL034402,
			AL117344,	U91325, AI		AC008101, AC004638
			Z86090, AC	AC004526, AC	AC002073, AI	AP000038, AP000106

				AC005740 Z95331, AP000194, AC002996, AL049869,
				ACO06556 AP000114 AP000046
				, iicoccoso, iicoccess, iiicoccos acoleoss, acoo4890, acoo402
	_			2010001 (05010001) (0501001) (0501001)
				, Arcocoas, Accouck
				, ACCOUNTE, ACCOUNCY, ACCOUNTS,
				, AF205588, AC00522
				AC002477, AC016830, Z94044, AC006146, AC004019,
				AC006077, AL117330, AL035089, AC009516,
				AP000036, AC006023, AC002400, AB000882,
				AC004020, AC004821, AC004814, AL132777,
				AL031311, AL117337, AC006064, L78810, AP000556,
				AC004699, Z84466, AC005332, AL109627, AL121653,
	-			Z93244, AC005969, AL022312, AC006958, AC005484,
				AL035455, AP000050, AL049635, AC003051,
				AC005488, AC006040, AC005562, AL133163,
				AC003029, AC004815, AC007637, AC005037,
				AC006160, AF196969, AC005585
2032 HWI	HWLDZ74	892558	Preferably excluded from the	AA337226, AI963222, AA336474, AI709289,
			present invention are one or more	AL079710, AI333306, AI095635, AI148461,
			polynucleotides comprising a	AA593438, AA460382, N99226, F35658, F28539,
			nucleotide sequence described by	AI674747, AI263147, AI689623, AI703331,
			the general formula of a-b, where a	AI304941, H46234, AA634465, AA336555, AA337527,
			is any integer between 1 to 969 of	AC004150, AC006024, AC006116, AC006539, U82672,
			SEQ ID NO:2032, b is an integer of	AC005592, AC007204, Z98747, AC006271, AC004045,
			15 to 983, where both a and b	AC007993, AF146191, Z54951, AC007284
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:2032, and where b is greater	
			than or equal to a + 14.	
2033 HPJ	HPJEB77	892563	Preferably excluded from the	H09290, AA806214, AA427513, AI904853, AA126879,
			present invention are one or more	AI910856, AW015950, AA134019, AA292157, AC009514
			polynucleotides comprising a	
			nucleotide sequence described by	
			the general formula of a-b, where a	
-			is any integer between 1 to 708 of	
			SEQ ID NO:2033, b is an integer of	

			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:2033, and where b is greater	
			than or equal to a + 14.	
2034	HNTST71	892820	Preferably excluded from the	W93943
			present invention are one or more	
			polynucleotides comprising a	
			nucleotide sequence described by	
	_		the general formula of a-b, where a	
			is any integer between 1 to 541 of	
			SEQ ID NO:2034, b is an integer of	
			15 to 555, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:2034, and where b is greater	
			equal to a + 14.	
2035	нсороэ2	893223	Preferably excluded from the	AA641005, AI762083, AI587618, AA143709,
			present invention are one or more	AW299688, AA524042, AI686577, AA143723,
			polynucleotides comprising a	AA534417, AW000937, AI924527, AI924182,
			nucleotide sequence described by	AA143746, AI478257, AW338896, AA999953,
			the general formula of a-b, where a	AI625051, AI417467, AA125991, AA233660,
			is any integer between 1 to 1070 of	AA233546, AA612904, AA826318, AI597567,
				AA906335, AA143761, AA126071, AI873680,
			15 to 1084, where both a and b	AI380837, AA056595, AA862082, AI910769,
			correspond to the positions of	AI380247, AA411502, AA328454, AI927431,
			nucleotide residues shown in SEQ ID	AA481473, AI368169, AA434336, AI002848,
			NO:2035, and where b is greater	AA056638, AW177469, AW177487, AI829000,
			than or equal to a + 14.	AA468833, U54603, AI916081, AW352026, AW365560,
				C00614, AW178439, AW292063, AW177675, AF216312,
				E13203
2036	HWLCU24	893457	Preferably excluded from the	AA479821, AA432116, AI571125, AW016789,
			present invention are one or more	
			polynucleotides comprising a	AA938157, AI422352, C06416, AI051837, AA425359,
			nucleotide sequence described by	W63640, AA479700, T66755, AW235659, AI978666,
			the general formula of a-b, where a	

	is any integer between 1 to 331 of	D52448, H49249, N54156, AA836066, AL043731,
	SEQ ID NO:2036, b is an integer of	
	15 to 345, where both a and b	A1954997, A1954988, A1589450, AA609914,
	correspond to the positions of	AI912009, AI218832, AI951761, AA609757, R77260,
	residue	R60869, AI460050, AW058594, AW300537, AA782792,
	NO:2036, and where b is greater	AA458911, N26791, AA708893, AI168124, W74653,
	equal to a + 14.	AI148331, AA188960, AI114875, AI915018,
		, T05685, AW
		AI745505, AA676964, H01261, AA129320, AA456251,
		AI653352, AA890006, AI096408, AW170047,
		AI247405, AI263393, AI081330, AI379150,
	_	AW015475, AA342341, H95038, AI814630, R08763,
		AI382384, AI273553, AI748817, N47474, AC005062,
		AF071240, AC005204, D37888, AF001893, AC005839,
		U46840, AC005082, AJ249224, X87116, D37887,
		Y09257, X96585, AL03
		AF175325, E15279, Z84484, AC005992, AC007298,
		3007917,
		AC007216, AC018769, AC009946, AL049543,
		AC005483, X79482, AJ388050, AC005884, Z93942,
		U09051, AC006112, AC002543, AF154112, AC004903,
		AF112374, AC006989, AF227510, AL109753,
		, Z83818, AE
		, AL022069,
		, AC004659
-		AC006151, X59370, Z83745, AC006196, AL078581,
		AC004001, X52617, A79336, U08407, AC005938,
	-	Z97180, AC004620, AC004533, AC006992, AP000459,
		AC002454, AC004849, AC006374, AL024506,
		AF178030, AL117338, AL109847, AC007320, M27933,
		AB017353, Z98043, AC005502, AL031177, Z96253,
		AF146793, AL049588, AF130342, AC012152,
		AF185591,
		35, I66426, AC003993, AL008723,
_		1185195

		_									_									_				_										
A13477,			AC005345,																L036525,							AI557378,								
AC007237,	AC053356,	AP001171,	U08869, A	AC005262,	AL023280,	AC005355,	AC007021,	AF015149,	AC007226,		AI064817,	AI174949,	AL037212,	AI133103,	AI133447,	AI708887,	AA618404,	AA468368,	AA149472, AL036525,	AA730806,	AI133076,	AA130534,	AI734894,	AL046874,	AA641711,	AA613948, A	AA188546,	AA876497,	AA211174,	AI978768,	AA937682,	AA583899,	AA610163,	AA151710.
AC016831,	AF109076,	AC005835,	AJ010688,	AC006478,	AF061032,	AC007058,	AC004391,	AF002166,	AF064863,		AI114520,	AI207400,	AI174746,	AA826080,	AL037712,	AIS57510,	AA639310,	AA176793,	C18264, AJ	AA533271,	AA155674,	AA176952,	AA293175,	AI954154,		AA657662, AA	AA149557,	AI031781,	AA493596,	AI986169,	AA180918,	AI453086,	AI133019,	AA469011,
AC004959,	AC000159,	AP000066,	AC011456,	AL023279,	AL079305,	AC005105,	AR031020,	AC005225,	AC006031,	U91328	AL037682,	AL037211,	AA661919,	AW131769,	AL047790,	AA196323,	AI557501,	AA528236,	AI720756,	AA723030,	AA524681,	AI735145,	AA115162,	AA888633,		C17903, AA	AI832615,	AI707485,	AI708877,	AI557052,	AI872466,	AA533010,	AI205258,	AI535649,
AE000658,	AL109623,	AC002041,	AC007970,	AC006972,	AP000884,	AL133241,	AC004108,	AC005513,	AC005723,	AR036572,	AI133205,	AL036965,	AI174789,	AI133183,	AI064872,	AI557213,	AI064799,	AA130931,	AA643792,	AA176099,	AA814574,	AA211175,	AA526147,	AA188082,	AI253288,	AA468936,	AA069837,	AA535388,	AW270369,	AA143743,	AI253289,	AA070665,	AA885561,	AI613175,
											7	present invention are one or more	polynucleotides comprising a	nucleotide sequence described by	the general formula of a-b, where a	is any integer between 1 to 1200 of	:2037, b is an intec	15 to 1214, where both a and b	d to the po	de residues s	NO:2037, and where b is greater	than or equal to a + 14.												
	_										893827							•																
											HSDJY15																							
											2037															_								

931, AA502034, 752, AI564738, 381, AI564738, 175, AA579454, 517, AI880251, 886, AI267882, 339, AA086434, 578, AA197115, 704, AA659697, AA659428, AA935460, AA935460, AA935460, AA935460, AA935460, AA935460, AA935460, AA749886, AI720230, AA081105, AA658333, 102, AA6583955, AA688903, 170, AM152114, 170, AM152114,	AA578931, AA502034, AA088752, AI564738, AI718381, AI057631, AA1569517, AA579454, AI569517, AA579452, AI523331, AA947056, AI523331, AA947056, AI620886, AN151535, AI889237, AA197115, AI889237, AA197115, AI801089, AA935460, AI630885, AI707630, AI630885, AI707630, AI630885, AI707630, AI630885, AI707630, AI630885, AI720483, AI637158, AI720483, AI637158, AI720483, AI832890, AA394073, AA618229, AA081105, AA618229, AA653867, AA618302, AI459667, AA618302, AI460107, AA618302, AI460107, AA506661, AA653864, AA50661, AA653867, AA618302, CI7416, AA AA50661, AA653867, AA132524, AI460107, AA226422, AA563955, AA192955, AI688903, AA6182955, AI688903,	AA578931, AA502034, AA088752, AI564738, AI718381, AI057631, AA879175, AA579454, AI569517, AI880251, AA100886, AI267882, AI523331, AA947056, AI523331, AA947056, AI889237, AA197115, AI889237, AA197115, AI889237, AA197115, AI801089, AA935460, AI630885, AI707630, AI67096, AA856914, AI832890, AA856914, AI697158, AI720483, AI697158, AI720483, AI697158, AI720483, AA618229, AA081105, AA618229, AA081105, AA618229, AA653867, AA618302, AI459667, AA618302, AI460107, AA618302, AI460107, AA226422, AA563955, AA226422, AA563955, AA192955, AI688903, AA6192955, AI688903,	AA088752, AI564738, AI718381, AA579454, AI718381, AI564738, AI569517, AI880251, AA579454, AI569517, AI880251, AA100886, AI267882, AI563331, AA947056, AI89237, AA197115, AI89237, AA197115, AI89237, AA197115, AI89237, AA659697, AI697158, AI720483, AI72038, AI72039, AA618229, AA61820, AA56661, AA658333, AA504102, AA658333, AA504102, AA658333, AA504102, AA658395, AI721188, AI720479, AA60898, AI832459, AA60898, AI832459, AA650170, AM52055, AI688903, AA650170, AM52055, AI688903, AA650170, AM52055, AI688903, AA650170, AM52055, AI688903, AA650170, AM520170,	10, AI45337	AA513214, AI812066,	AA552282, AI862343,	AI133109, AI625924,	AA522574, AA130876,	AA074099, AI242732,	AI799288, AI041459,	AL047605, AA101240,	AI921645, AI735153,	AI719836, AI610718,	7010, AA46	AI749770, AI635150,	AI147985, AA652921,	AA536131, AI269472,	AA533389, AA602791,	AI253340, AI801192,	AI217009, AA603147,	AA661870, AI091584,	_		08, AA57584	, AI21620	AA46921	AA58668	29,	٦,	AA771977, AA526043, AI720323,	AI080487, AI720329,	A183	AI366465, AI459785,	, AA09	AA708210, AA485747,	AA174120, AW166854,	C18862, AA775370, AI031761,	AA548147, AA545759,
				, AA502034,	, AI564738,	, AI057631,	, AA579454,	, AI880251,	6, AI267882,	, AA947056,	99, AA086434,	8, AW151535,	AA197115,	, AA669697,	9, AA935460,	, AA856914,	5, AI707630,	AA659428,	539, AI273169,	8, AI749886,	3, AI720483,	0, AA394073,	9, AA081105,	3, AI557420,	2, AI459667,	9, AI720230,	6, AA744189,	1, AA658333,	2, AA603867	2, C17416,	7, AI469695,	0, AI832984,	4, AI460107,	2, AA563955,	8, AI720479,	AI832459,	5, AI688903,	, AW152114,
	·			AA578	AA088	AI718	AA879	AIS69	AA100		AI499	AI926	AI889	AI832	AI801	AI670	AI630	AI475	AI124	AI720	AI697	AI832	AA618	AA566	AA829	AL047	AA468	AA506	AA074	AA618	AA502	AI721	AI832	AA226	. AA211	AA600	AA192	AA650

2038	HSAAR81	893842	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 442 of SEQ ID NO:2038, b is an integer of 15 to 456, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:2038, and where b is greater than or equal to a + 14. Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 580 of SEQ ID NO:2039, b is an integer of 15 to 594, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:2039, and where b is greater than or equal to a + 14.	ANIS66365, AIS72029, V X93334, U09500, X93333 D38113, X93347, D381115 , AJ010580, AJ010582, X13302, X13302, X13305, X134861, AA315762, AA974853, AA631397, AIA60867, AI983974, W6 AMC5199, AA327340, AA327154, AI350070, AA384635, W81242, AA327340, AA327154, AI187009, AI832569, AI832
				AII83838, AA468623, AA928702, AW176584, AI973212, AI749833, T29881, D25724, AA314975, AA029584, W95644, AA574221, I95749, I15203.
2040	HCNSE58	893867	Preferably excluded from the	25654, U25656 AI143630, AA315762, AA5524

			present ingention are one or more	AA974853	AW167728.	AA716097. AI332337.
			leotides comprising a		AA953818,	
			nucleotide sequence described by		AA580138,	AA507878, AA614535,
			the general formula of a-b, where a	AA552321,	AI762067,	W32686,
			is any integer between 1 to 639 of	W60304, AI	AI983974, AI973218,	AA633399, AI98
•			SEQ ID NO:2040, b is an integer of	AA554791,	AA314975, N74131,	
			15 to 653, where both a and b	AA337636,	W81242, AJ	AI832569, W81706, AI183698,
			correspond to the positions of	AA468623,	AI459674,	AI749833, AI460270,
			nucleotide residues shown in SEQ ID	AA928702,	AI187009,	AW364159, W95642, T29881,
			NO:2040, and where b is greater	AA345906,	AI474125,	_
			than or equal to a + 14.	AA384635,	AA384262,	AA327250, AA336734,
				AI561269,	AA327500,	AA327546, AA574221,
				AA327340,	AA029584,	AA327502, AI699171,
				AW176584,	AA327154,	AA532852, AW188590,
				AA558976,	AI560870,	AI749877, AA319354,
•				AW007096,	W95643, AJ	W95643, AA337338, AA384655, AA029583,
_				W95644, AW392670,		AW291863, Z99396, AL119319,
				AL037205,		
					AL119439,	
					AL119522,	AL119457, U46347, U46351,
				AL119483,	AL119418,	AL119418, L15203, I95749, L08044,
				72	5656, U25	5654, AR060234, AR066494,
				A81671, AB	AB026436	
2041	HSVCD79	894012	Preferably excluded from the	AA429308,	AW138602,	AW024259, AA558588,
			present invention are one or more	AI492469,	AI367813,	AA428240, AA719541,
			polynucleotides comprising a	AA888930,	AI190902,	C14850, AI217028, D60222,
			nucleotide sequence described by	AI286160,	AA737138,	R79200, H64703, R79465,
			the general formula of a-b, where a	AA737139,	AI268290,	AF023259
			is any integer between 1 to 1902 of			
			SEQ ID NO:2041, b is an integer of			
			15 to 1916, where both a and b			
		_	correspond to the positions of			
			nucleotide residues shown in SEQ ID			
			NO:2041, and where b is greater			
2042	HSIFA27	894051	Preferably excluded from the	AI972556,	AI968208,	AW274901, AI744720,

			present invention are one or more	AI885290, AA449113, AW152432, AI479938,
			polynucleotides comprising a	AI800087, AW390446, AI800088, AI799502,
			nucleotide sequence described by	AI859002, AI423145, AW088405, AI858842,
•			the general formula of a-b, where a	A1990019, A1809596, A1401062, A1360174,
			is any integer between 1 to 1581 of	AW197421, AI689608, AW197663, AW103934, N42254,
			SEQ ID NO:2042, b is an integer of	AI218225, AI206902, AI376613, AI219568, N59385,
			15 to 1595, where both a and b	AA053930, AA534904, AI656541, AI128371,
			correspond to the positions of	AI360254, AI285163, N32810, AA428038, N39444,
			nucleotide residues shown in SEQ ID	AA776360, AW088291, AI817703, AA421739,
			NO:2042, and where b is greater	AI565066, AI674914, AW190558, AW194393,
			than or equal to a + 14.	AW276699, AI361508, AI824832, AW451191, R91784,
	***			AW390451, AA427924, AA257059, AW071546,
				AI081359, AI189019, AI002857, W93989, AW206484,
				H55900, AA034237, AA127466, AW188281, AI290045,
				AA447735, AW027775, AA773930, AI633932,
				AA327290,
				W93800, AI690373, AW027793, AI143661, R59973,
				AA503464, R82261, R91785, N63596, AW276891,
				AW276821, AW182096, H01166, H01251, N29781,
	-			H24046, H13082, R27203, AI811525, AA055340,
	-	•		AA319583, AA358644, AA904821, AI274485, R27202,
	•			9, R46792, N57202, R67153,
				H61875, AI783927, AA453668, C15384, AB018305
2043	HTTKV46	894121	Preferably excluded from the	AI678077, AI884863, AI869333, AI884942,
			present invention are one or more	AI859296, AA829937, AW250313, AW300936,
			polynucleotides comprising a	AI571293, AW273060, AW248281, AA582906,
			nucleotide sequence described by	AA928110, AA283711, AI589898, AI038859,
			the general formula of a-b, where a	AA594105, AA828316, AA906924, AA938955,
			is any integer between 1 to 1047 of	AW170665, AW172642, AW248955, AA975490,
			SEQ ID NO:2043, b is an integer of	AI123879, AI367867, AI826097, AW272915,
			15 to 1061, where both a and b	AW070748, AA316879, AI089508, AI086474,
			correspond to the positions of	AA661759, AI566244, AI015067, AI538087,
			nucleotide residues shown in SEQ ID	AW245061, AW000868, AW409921, AA688299,
			NO:2043, and where b is greater	AW250988, AA827720, W58033, AI953468, AA211097,

than or equal to a + 14.	AW078745, AI891144, AA994072, W79220, AI471577,
	W74508, AI922589, AW102638, AA918328, AA826730,
	3, D56355,
	5, AW340401
	, AA290724, T19021,
	AI307442,
	AW075100, AA380031, M91218, AW073433, AI802854,
	, AW071289, AI349002, AW07517
	AI307208, AW072721, AI334909, AI312145,
	_
	AI307478, AI348921, AI252839, AI307493,
	AI255068, AW073456, AW072496, AW302738,
	, AW301481,
	, AI345565, AI33488
	, AI25246
	, AI255052,
	, AA824526,
	, AI309390, AI3
	AI340619, AI252075, AI254764, AI251262,
	3, AW302733, AW073049, AI25123
	7, AI247038, AW072901,
	, AW271039,
	5, AI269525,
	AI340643, AI054057,
	2, AW074866, AW302327,
	AI054172, AI053900,
	, AI271496,
	AI252427, AA993616, AI307473, AA496372,
	AA464729, AI566787, AI885746, AA496649, T90849,
	н
	AW249375,
	AI345130, AI254134, AI340511, AI349742,
	AI334895, AI307507, AI310927, AI336488,

			NO:2045, and where b is greater than or equal to a + 14.	
2046	HWLVS05	894631	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 1425 of SEO ID NO:2046. b is an integer of	AI952147, AA827782, AI523970, AW008938, AA236865, AI673370, AW043829, AI143323, N36986, AA306716, AI361743, AA460666, AW080829, AI914077, AI214786, AA862831, AI963652, AI913070, AI805253, AI423188, AI003936, AA994686, AA130868, AA533231, AI358965, AI873692, AA569719, AA865951, AA644481,
				ma mm ww
				A57389, AL137562, AF158248, U72071, X79812, AL049959, AF070632, U92068, AJ131955, AF169154, AF030165, Z30970, AL0956709, Z49258, AC006561, AL022396, Z98049, AC007370, AL049540, AL021391, U94316, AP000250, AP000133, AP000211, AP000030, AF162270, I80845, AF107018, U77594, AL080074, AR029580
2047	HCRMV27	894806	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 572 of SEQ ID NO:2047, b is an integer of 15 to 586, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:2047, and where b is greater	AL134920, AL042896, AL119443, AL042965, U46341, AI142139, AL119418, U51899, A81671

2000	LCDOID	110100	Destaurable seement from the	
7040	DCROI22	110460	_	
			present invention are one or more	AL134531, AW372827, AL119439, AL119484,
			polynucleotides comprising a	AL119363, AL119391, AL134528, AL119444,
			nucleotide sequence described by	AL119418,
			the general formula of a-b, where a	A81671
			is any integer between 1 to 881 of	
			SEQ ID NO:2048, b is an integer of	
			15 to 895, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:2048, and where b is greater	
			than or equal to a + 14.	
2049	HCQAF06	894818	Preferably excluded from the	
			present invention are one or more	
-	_		polynucleotides comprising a	
			nucleotide sequence described by	
			the general formula of a-b, where a	
			is any integer between 1 to 129 of	
			SEQ ID NO:2049, b is an integer of	
			15 to 143, where both a and b	
			correspond to the positions of	
			NO:2049, and where b is greater	
			than or equal to a + 14.	
2050	HKCSA83	894820	Preferably excluded from the	AW360811, AW177440, T03269, AW375405, AW178893
			present invention are one or more	, AW367950
			polynucleotides comprising a	AW179328, T48593, AW178906, AW375406, D80439,
			nucleotide sequence described by	AW377672,
			is any integer between 1 to 562 of	D80193, D59927,
			SEQ ID NO:2050, b is an integer of	
			15 to 576, where both a and b	D59275, AW178762,
			correspond to the positions of	
			·O	D80227, D59502, AA305409, AW378532, AA305578,
			50, and w	AW377676, AW352170, AW178907, AW178908, D80251
			than or equal to a + 14.	AW178914, C06015, AW378533, D45260, AI525923,

				C03092, AA285331, AW378542, AI525917, AA809122,
				Fise4', Aisesse0, Amsifie4, 3, AIS25925, IS0126, IS0132,
				AR066487, AR060138, A84916, A45456, A67220,
				A62298,
				A43192, AR008278, X67155, Y12724, A43190,
				AR038669, AF058696, A25909, AR008443, AB002449,
				D88547, Y17187, D50010, A63261, A70867, X82626,
				AR062872, AR025207, AR008408, AR016691,
				AR016690, U46128, A64136, A68321, D13509,
				I14842, AR054175, AR060133, X68127
2051 HS	HSBA104	894824	Preferably excluded from the	-
			present invention are one or more	C14389, D58283, D80188, D80391, D59787, D51423,
			polynucleotides comprising a	AW360811, D80247, D50979, D80196, D80439,
			nucleotide sequence described by	D80522, C14014, D80212, D51022, D59859, D80022,
			the general formula of a-b, where a	D80166, D80195, D59467,
			is any integer between 1 to 566 of	D80164,
			SEQ ID NO:2051, b is an integer of	D59502,
			15 to 580, where both a and b	D80248, D81026, D80269, D80366, D80219,
			correspond to the positions of	, C15076
			de	D59889, D80193, D80133, D80045, D80024,
			NO:2051, and where b is greater	AA514186, AA514188, D80302, D80157, D80378,
			than or equal to a + 14.	AW177440, D51759,
				D80251, AW178893, T03269, C06015, AW377671,
				AW375405, H67854, AA809122, AW178906, AW366296,
	-			AW360817, AW179328, T48593, AW375406, AW378534,
•				F13647, AW179332, AW377672, AW179023, AW178905,
				AW177731, AW378528, AW178762, AW179019,
				AW378532, AW352170, AI525923, T11417, C03092,
				H67866, AW179020, AW377676, AW352171, AI525917,
	· · · ·			AW178907, AW178908, AW179024, D51250, C14227,
				, D58101,
				D59317, AW367950, AW177456, T03116, AI525227,

		178980, D58246, AW178986, D81111,
		AW178774, D80258, AW179018,
		D80064, C14344, AW378533, AIS2
		AI525912, AA514184,
		, AW178911,
		163, D59551, D52291, AI
		D59627, AI
		8, AI557751, AI525222, D51213,
		AI525237
		T03048, D45273, C05763, T02974, Z21582,
		5, AI5259
		T0
		C13958, AI525238, T11191, D31458, AI525913,
		AC000047, AR008278, AB028859, AJ132110, A84916,
		A62300, A62298, A82595, AR060385, AR018138,
		AF058696, AB002449, I50126, I50132, I50128,
		Y17188, AR016514, X67155,
		A94995,
		Y12724, A43190, AR038669,
		8, Y09669
		D34614,
		Y17187, AR008277, AR008281,
		D50010, A
		X82626, AR008408, AR016691, AR016690, U46128,
		207, X64588,
+		9, X68127,
HCQCD80 894827	<u>.</u>	4, D80024, D51079,
	present invention are one or more	1, D80366
	otides comp	9, AA809122, D51053, D80248
	eotide sequence	C14389, C14014, D80268, D80439, D58246, D45273,
	eral formula of a-b, wher	0, D45260,
	teger between 1 to 557	6, C15076, D80166,
	:2052, b is an int	D80210, D51799, D59551, D80240,
	where t	D80219, D58283, D8
	d to the positions o	AA305409, D81026, D80269, D80022, C14331,
	nucleotide residues shown in SEQ ID	D80195, AA305578, D59627, C14973, Z33452.

	T03269, AW377671, AW375405, D59653, C75259
	4, AW366296, AW178906, AW360844, AW3
	8, T48593, AW375406, D59373, AW378534,
	AW377672, AW179023
	AW178762,
	2, AI525923,
	AI535686, D80064,
	N66429, AW179020,
	D80258, AW352170
	AW179024, D58246,
	D51250, AW360841, AW
-	AW177505, AW176467,
	AW178909, AW179004,
	7,
	4, C14227,
	ΑI
	51221, D59474, D60010, AI535959,
	AIS57774, D60214, AA514184, AW179009, AW179012,
	AW178911, AI525227, AW378543, AW378525,
	2, AW352163, D52291,
	AW177734, C14957, AIS2
	D80949, D59627, AI525242,
	, AI525215, D51213,
	, D59976, A
	, AB005289, AF0787
	A82595,
	AR018138, A62300, AR06
	, AJ132110
	, IS0126,
	77, X82626,
	AR060138,
	, A26615, AR052274,
	, AR038669, A25909,
	, AR066487, X68127,
	AR008443,
	AR016690, D50010, D88547, A63261, A70867,

				AR062872, A68321, D. AR032065	AR062872, AR008408, A68321, D13509, AR06 AR032065	AR025207, 50133, 1824	AR062872, AR008408, AR025207, I79511, A64136 A68321, D13509, AR060133, I82448, AF123263, AR032065	
2054	нсоре22	894831	Preferably excluded from the present invention are one or more	N58518, A	AA699859, AA	AA677543, AC	AC006556	
			nucleoride sequence described by the general formula of a-b, where a					
			is any integer between 1 to 829 of					
			SEQ ID NO:2054, b is an integer of					
			15 to 843, where both a and b					
			correspond to the positions of					
			nucleotide residues shown in SEQ ID	*				
			NO:2054, and where b is greater					
			than or equal to a + 14.					
2055	HWLVU33	894832	Preferably excluded from the	AA775419,	AI273235,	AI754154,	AI446402,	
			present invention are one or more	AI640735,	AI468600,	AA602645,	AI675266,	
			polynucleotides comprising a	AA460180,	AI382693,	R40043, AA813916,	1813916, AL119457	57,
			nucleotide sequence described by	AI110849,	AL042544,	AL042382,	AL119399,	
			the general formula of a-b, where a	AL079794,	AA225339,	AI468872,	AW262565,	
			is any integer between 1 to 739 of	AI499463,	AI680498,	AI362637,	AI491852,	
			SEQ ID NO:2055, b is an integer of	AL043326,	AL135661,	AI224992,	AI648684,	
			15 to 753, where both a and b	AL045903,	AI679990,	AI590118,	AW148716,	
			correspond to the positions of	AI446092,	AI282326,	AIS54245,	AI857296,	
			nucleotide residues shown in SEQ ID	AI701074,	AI569616,	AI446605,	AW087445,	
			NO:2055, and where b is greater	AW071417,	AI636445,	AI567360,	AI591316, N42321	121,
			than or equal to a + 14.	AI269696,	AI500039,	AI758437,	AI612920,	
				AIS70384,	AI801766,	AA640779,	AA287231,	
				AI811344,	AI520785,	AI886124,	AI690312,	•
				AI590120,	AI475451,	AL040243,	AI554427,	
				AI273142,	AI097248,	AW103893,	AW150578,	
				AI869367,	AI868831,	AI633419,	AI433976,	
				AI280747,	AW302988,	AW274192,	AI687065,	
				AI612759,	AI800453,	AI800433,	AI684265,	
				AW023590,	AI273048,	AI539771,	AI816947,	-
				AI610756,	AI274013,	AI500146,	AI537677,	

951	0	(Ω)	3915	
AW075084,	AW118512,	AW131954,	AW196141,	
. 9	8597	155	168	
28177	52813	5332	9857	
002	455	0243	34305	
AI799199,	3204	AW102785,	5129	
64	AI349933,	40	88895	
 AW088903,	AL038565,	AW088793,	AI866002,	
 AI828731,	AI866608,	AI866111,	AI919345,	
 AW162071,	AI251830,	AI366549,	AI636719,	
AW238730,	AI802542,	073	AL036214,	
AW074993,	AI349614,	AI800411,	AI538085,	
445	AW268253,	986	AI312152,	
AI952360,	AI264741,	AI340582,	AI784252,	
AW132034,	AW193000,	AI349937,	AI702406,	
AI567993,	AW301410,	AI571909,	AI349004,	
AI620287,	AI917055,	AI307708,	AI318280,	Ī
 38	AI308035,	AL036146,	AL036759,	
85	950	370	AI923768,	
AI682743,	AI678302,	AL079963,	AW403717,	
AW071349,	947	AW268220,	AIS60099,	
AW103371,	AI273843,	AI521012,	AI270707,	
049	AI281837,	82	AI801152,	
4	AI632033,	AI434223,	AL079741,	
\sim	AL045500,	AI922901,	AI249257,	
909	AI569583,	AI572787,	AI564247,	
 AI282281,	AW075351,	AI925156,	AW169653,	
	AW148320,	AI608936,	AL119863,	
AW075413,	AIS00077,	AW167410,	AI282903,	
900	AI862144,	AI439717,	AI567612,	
AI284131,	AI570989,	AI312428,	AI619749,	
AI567351,	25	AI431424,	AI250663,	
982	17	on	AL036980,	
 3838	355	O)	AI572676,	
AL036802,	AI269862,	AW071177,	AI476109,	

AI345735, AI648663, AL036396, AI950664,
AI872074, AW168723, AI538716, AA613907,
AW089572, AI334884, AI348897, AL036274,
AI612885,
AI340627, AI634224, AI445237, AW151138, Y11587,
AF158248, S68736, I48979, AL122093, AL122050,
AF125949, AL133640, AL110196, AL133016,
AL117457, AL137557, I48978, A08916, AF113013,
 I89947, AF078844, A08913, S78214, I89931,
A93016, AL080137, AF118064, L31396, L31397,
AF017152, AL080060, AF113694, AF113691,
AL050393, AL133565, AF113690, AF113019,
AF090934, A65341, AL137459, AL137527, U42766,
AF090943, AF090900, AF113676, AF111851, I49625,
AL133557, AL110221, AF125948, AL050146,
AB019565, E03348, AF118070, X84990, AL050149,
AL133606, AF104032, AL049314, AL133093,
 AL080124, AL049452, AF113677, Y11254, AL050116,
AF091084, U91329, AL122121, AF017437, AL049938,
X63574, AL
AF090903, AL096744, AF146568, AF090896,
AF090901, AR059958, AF079765, AF106862, Y16645,
AL117585, AL133075, X82434, AJ000937, AL133080,
 AF113699, AL049466, AL137550, AL050277, E07361,
AL049382,
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L117435, A58524,
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A08912, I33392,
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061943,
AL137463, X93495, U35846,
.03736, AL122110, U80742, U726
AL137560, I09360, AL049283, AJ012755, AF119337,

			AF087943, AL122111, AF067728, AL080159, AL133072, X98834, AL050172, AL110197, E08263
			69,
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			AL137476, AF026124, U96683,
			Y09972, I26207, AL133077, AF111112, AF057300,
			AF057299, M30514, AL137556, AF132676, AF061836,
			A08911, AL137523, I00734, I66342, U68387,
			AL133104, AL080074, AR013797, AL137526,
			AL133098
			AF067790, A45787, E05822, AL110280, AF106827,
			3006371, AL133067, AF153205,
			AF162270, AF081195, L19437, AL117440, AR038854,
			AL137533, A90832, Y07905, X62580, AJ006417,
			X53587, AF111849, L30117, AF061573, U49908,
			AL137705, AF008439, AC004200, X87582, U58996,
			1, E08631, AL
			AR054984, AL080158, AL023657
2056 HAJAY88	894842	Preferably excluded from the	AI623302, AI432655,
		present invention are one or more	AI431337, AI431328,
		polynucleotides comprising a	AI432666, AW081103,
		nucleotide sequence described by	AW128900,
		. formula of a-b, where	AI431346, AI432662,
		teger between 1 to 4002	AI431255, AI431243,
		SEQ ID NO:2056, b is an integer of	
		15 to 4016, where both a and b	, AI432665,
		correspond to the positions of	AI431351, AI431345, AI432672, AI431254,
		nucleotide residues shown in SEQ ID	AI432676, AI431241, AI432673, AI432658,
		NO:2056, and where b is greater	AI432674, AI431340, AI432664, AI431307,
		than or equal to a + 14.	AI431316, AI791349, AW128897, AI432657,
			2520, AW129223,
			AI432643, AI431751, AI492509, AI492510, Y17793,
			AF064854, AF019249

2057	HCRPM46	894878	Preferably excluded from the	AL119319, AW392670, AL119418, AL042551,
			present invention are one or more	AW372827, AW363220, AW384394, AL119497, Z99396,
			polynucleotides comprising a	U46341, AL119483, AL119457, AL119443, AL119324,
			nucleotide sequence described by	AL119484, AL119363, AL119341, AL119391,
			the general formula of a-b, where a	AL119355, AL134531, AL134518, U46351, U46349,
			is any integer between 1 to 573 of	AL042965, AL119399, AL119335, AL119522,
			SEQ ID NO:2057, b is an integer of	AL119396, U46350, U46347, AL119496, AL119444,
				U46346, AL134528, AL042975, AL134538, AL042542,
			correspond to the positions of	AL037205, AL134920, AL134533, AL119439,
			nucleotide residues shown in SEQ ID	AL042614, U46345, AL043019, AL042984, AL043029,
			NO:2057, and where b is greater	AL042896, AL043011, AL042970, AL042450,
		_	than or equal to a + 14.	AL042544, AL043003, AL119488, AL119464, A81671,
				AR060234, AR066494, AB026436, AR054110, AR069079
2058	HOE0Q19	895122	Preferably excluded from the	AA307684, AA232750, AI417539, AA100160,
			present invention are one or more	AA864846,
			polynucleotides comprising a	AW364482, AW364479, AR044133, AR044123, AR044135
			nucleotide sequence described by	
			the general formula of a-b, where a	
			is any integer between 1 to 1049 of	
			SEQ ID NO:2058, b is an integer of	
			15 to 1063, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:2058, and where b is greater	
			than or equal to a + 14.	
2059	HKGBP52	895303	Preferably excluded from the	AW058657, AA400627, AI692280, AI342528,
			present invention are one or more	AI743405, AA400382, AI675621, AI808100,
			polynucleotides comprising a	
			nucleotide sequence described by	AW135173, AI343951, AI299820, AA393033, T03738,
			the general formula of a-b, where a	N24268, H98701, AI040531, R56558, H54669,
			teger between	AI830628, H01460, C16675, AA707616, H00353,
			SEQ ID NO:2059, b is an integer of	AI146912, H01555, R21829, AI755214, AI754567,
			15 to 2716, where both a and b	\mathbf{H}
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	AI569100, AI858691, AI583142, AW192599,
			NO:2059, and where b is greater	AI077941, AA176978, AA704393, AA602906, H00307,

	than or equal to a + 14.	AA491767, AA719073, AA659832, AW270385,
		AI884383, AI354423, AI061313, AI590458,
		AI679002, AW270255, AI679759, AI926728,
		AI590499, AW069227, AI732502, AI791458,
		AI609972, AI754336, AI590580, AI499376,
		AW022934, AI753113, AW277253, AW438856,
.		AA584765, AA484892, AI791659, N71685, AA444166,
		H85383, AA171892, AW089950, AI572680, AA715173,
		AI636734, AA720702, T57096, AI707788, AA622801,
<u>-</u>		T71936, AI431513, AA583386, AA525753, AI753488,
		AL080317,
		L035455, AJ010770,
	-	7, AC007216, AL035454
		AF129756, AC00658
	-	
_		AC005088, AC005280, AC010170, AC004685, Z82976,
		AC006511, AC004148, AF045555, AC002551,
		_
		, AC006241, AL109628
		AC005365, AL109759, AL023575, AL049759,
		, AC005231,
		1, AP000135
		9, Z81364, AF176915, Z
		, AL034417, AL009181
		6, AL109798,
		7
		AF134726, AF030453, AC005488, AC006141,
		AF024533, AC007055, AC002990, AC006930,
		AP000512, AC007250, AC007687, AC004534,
		AC007308, AC005332, AL034429, AL021331,
		AC004518,
		., AC002395, AL080243,
		AC004922,
		M63544, AC010077, AB023048, U91319, AC004895,
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			AA448779, AA362005, AA383368, C16419, F	AA448779, D57975, AI474663, AA627283, AW3516 AA362005, T06370, AA581145, F11298, H03672, AA383368, F08958, D62803, H03671, AI264956, C16419, F10353, AW388337, AA243374, AI79664	AI474663, AA627283, AW351677, AA581145, F11298, H03672, D62803, H03671, AI264956, J388337, AA243374, AI796664,
			AI /38552, AI539480,	A1695343, AW391667, A1800690 H87103, AW150643, AC008498,	5, AW39166/, AI800690, AW150643, AC008498, AL021997
HDPPB40	895675	امرا	AI223386,		
		present invention are one or more	AL043887,	ò	5, AI223392,
		polynucleotides comprising a	AI750397,	AA813783, AI911812,	2, AA253429,
		nucleotide sequence described by	AI799380,	F09731, AL043886, T81826,	T81826, AI221738,
-			T65287, T	T65235, AR052513, D50419	50419
		is any integer between 1 to 2581 of			
		SEQ ID NO:2061, b is an integer of			
		15 to 2595, where both a and b			
		correspond to the positions of			
_		nucleotide residues shown in SEQ ID			
		NO:2061, and where b is greater			
		than or equal to a + 14.			
HWL0129	895781	Preferably excluded from the	AC006050		
-		present invention are one or more			
		polynucleotides comprising a			
		nucleotide sequence described by			
		the general formula of a-b, where a			
-		is any integer between 1 to 540 of			
	_	SEQ ID NO:2062, b is an integer of			
	_	15 to 554, where both a and b			
-		correspond to the positions of			
		NO:2062, and where b is greater			
		than or equal to a + 14.			
HCRMJ47	895927	Preferably excluded from the	AW084003,	AA570505, AA526186	6, AW006250,
		present invention are one or more	AW007762,	AI458032, AA149494	4, AI799666,
		polynucleotides comprising a	AI341557,	AI084783, AI190971,	1, AI377966,
		nucleotide sequence described by	AI085276,		0, AW148913,
		the general formula of a-b, where a	AI380460,	AI123203, AI122890	0, AW007426,
		is any integer between 1 to 1834 of	AI863238,	AA603986, AI307748	B, AI921067,

			SEQ ID NO:2063, b is an integer of 15 to 1848, where both a and b	AA149490, W73595, AV	AA149490, AI280975, AI336463, W73495, W73595, AW149089, AI814701, AI766921,	AI336463,	1	AI367500, AW450642,
			correspond to the positions of	AA235464,	AI189309,	AW072576,	AI129064,	
			nucleotide residues shown in SEQ ID	AAS74230,	AA292528,	AA650188,	AI589229,	
			NO:2063, and where b is greater	AW294024,	AI580733,	AA037024,	AI288103,	
			than or equal to a + 14.	AA877009,	AI660255,	F24537, AA	578293,	AA047125,
				AA864573,	AI274628,	AW188597,	AI572782,	
				AA374109,	AI866359,	AA558228,	AA621604,	
				AI264439,	AA658397,	AI652870,	AA573559,	
				AA573997,	AI567038,	Z39737, AV		AW243333,
				T81066, A.	T81066, AI684973, AA034505,		AW377101, AA	AA372354,
		-		AA047126,	AB027466,	AR035961, AR037874	AR037874,	
				AR035966,	AR035967			
2064	HLDXE66	800968	Preferably excluded from the	AI500518,	AW328444,	AW327862,	AI971783,	
			present invention are one or more	AW328440,	AW328380,	AW328614,	AW327796,	
			polynucleotides comprising a	AW007896,	AI628924,	AW410322,	AW409642,	· <u> </u>
			nucleotide sequence described by	AW328007,	AW328376,	AW087373,	AI754439,	
			the general formula of a-b, where a	AW409590,	AI287514,	AA551550,	AA501684,	
			is any integer between 1 to 473 of	AI440000,	AA603360,	AI818460,	AI201181,	
			SEQ ID NO:2064, b is an integer of	AI610070,	AW409683,	AA535393,	AI699829,	
			15 to 487, where both a and b	AIS59540,	AI560651,	AA574413,	AI827247,	
			correspond to the positions of	AW328350,	AW328609,	AA513486,	AI754460,	
			nucleotide residues shown in SEQ ID	AW134985,	AI755116,	AW007719,	AA283266,	
			NO:2064, and where b is greater	AA854768,	AI497632,	AA772414,	AA496883,	
			than or equal to a + 14.	AW328320,	AA679713,	AI050044,	AW020501,	-
				AI619744,	AI339813,	AA886011,	AW250421,	
				AI274211,	F32918, AA	4579416, AA632536,	4632536, AI	AI567937,
				AI831479,	AI151481,	AI186976,	AA877933,	
				AI185119,	AI620681,	AA757769,	AI690593,	
				AA714364,	AA558105,	AI922235,	AA632723,	
				AA843775,	AI924171,	AI961721,	AW250755,	
				AW090148,	AA491636,	AI338728,	AI123375,	
	-			AW090155,	AW081336,	AI539209,	AI890302,	
			•	AW245791,	AI573062,	AI028444,	AI863898,	N64026,
				AI439763,	AA847963,	AI749978,	AW261931,	
				AI634383,	AI191638,	AI198771,	AI719450,	

	AI344453,	AI718439, AI26867	7, AI631303,
	AI253560,	AW250772, AI796657	7, AA536044,
	AA569292,	AW169077, AI570813	, AI69747
		AI355377, AA61027!	5, AI674831,
	AI114866, 1	AI565047, AI193415	, AI57145
			6, AA536172,
		AI660181, AW273029	9, AI818029,
	ر. د	AA580796, AI719806	6, W73177, AW080272
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			0, N91538, AW090784
	_	6, AI66015	ر م
	_	AI859783, AW028278	8, AA598891,
_	AI281231, 1	AI289421, AW30519	5, AA908802,
	AW073669, 1	AI800405, AI342580	0, AI432916,
	_		AI27285
	AI963461, 7	AI620289, AI523503,	3, AI891159,
	AI475307, I	R02544, F24388, N	F24388, N32326, AI270199,
	_	AW242012, AA448266,	6, AI557537,
	_	H	AA776791, AI206373,
	7,	AI510744, AA723534	4, AI831263,
	1,		2, AW118551,
	ς,	_	2, AA737215,
***	_	_	8, AW023162,
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	e,		_
-	7	_	8, N31753, AI583997,
-		AI333494, AA580751	1, AA879000, W85708
	AA508174, A	AW328608, AI439940	0, AI689023,
	٠	AI924195, AW188874	1, AA491865,
	AA312014, P	AA908266, AI160628	3, AI860497,
		AW084818, AI557538,	3, AW273989,
	7	AI735229, R16758,	R16758, AI333611, AW079820,
	_	.939, AW337470, AJ	F21939, AW337470, AI160685, AA507934,
	/KK 30000W	אטטטארמאל נטארט	CONTRACTOR TO CONTRACTOR CONTRACT

				AI969498,	AW245433,	M36072, AC	AC000089, X06705,	
				AJ224080,	AC004217,		X52138, AC002107,	-
				AL034417,	AB023058,	AP000521	AL022723,	
				AF055066,	AJ224082,	AC004192, AC004172,	AC004172,	
				AJ224081,	X15013, A	AJ224081, X15013, AC000399, AC005042,	3005042, Z84469,	
				D63790, AC004129,	2004129, AJ	L031736, AC	AL031736, AC007110, AL078595	
				AC002452,	Y17212, T	51109, TSST	AC002452, Y17212, T51109, T55719, T56886,	_
					T59899, T59	990, H50847	T59990, H50847, H98782, N24572,	
					N95637, W69'	735, AA025	W69735, AA025830, AA070711,	
					AA084650,	AA085276, AA102516	AA102516,	
				AA148893,	AA150738,	AA156887,	AA181948,	
				AA187531,	AA425933,	AA428802,	AA226324,	
				AA279495,	AA480450,	AA484692,	AA523996,	
				AA535068,	AA554440,	F15687, AA586409,	1586409, AA602157,	_
				AA603678,	AA610650,	AA632560,	AA580635,	
				AA730447,	AA737209,	AA862929,	AA863478,	
				AA885536,	AA886913,	AA954603,	AA962430,	
				AA975386,	AA976970,	AA991428,	AA999672, N87911	_
				AA641479,	AA129690,	AA211080,	AA400765, F20644	
				AA775513,	AA283334,	AI078081,	AI078082, T11296	_
				AA693434				
2065	HAIBM54	897234	Preferably excluded from the	AW245845,	AW245888,	AW247437,	AA226733,	
			present invention are one or more	AA019081,	AA325881,	AW247424,	AA324707,	
			polynucleotides comprising a	AI802708,	AA315689,	J04469, Z	J04469, Z13969, X59737,	
			nucleotide sequence described by	213968				
			al formula of a-b, when					
			teger between 1 to 561					
			:2065, b is an					
			15 to 575, where both a and b					
			correspond to the positions of					
			nucleotide residues shown in SEQ ID		•			
			NO:2065, and where b is greater					
			than or equal to a + 14.					
9907	HSXAX45	897524	Preferably excluded from the	AI459464,	AA808743,	AI144559,	AA861434,	
			present invention are one or more	AA404217,	AA630335,	AI831253,	AI248728,	
			polynucleotides comprising a	AI870869,	AA618605,	AI458793,	AI027413,	

 eotide sequence	AA918131, AI128366, AW405777, AI800139,
al formula of a-b, wher	AI805659, AA569324, AI138987, AI333605,
teger between 1 to 772	AA461611, AW189901, AA461439, AA586689,
 :2066, b is an inte	AA915895, AA991975, AA642111, AI033160,
 15 to 786, where both a and b	AA459952, AA503924, AA622287, AI126939,
 correspond to the positions of	AA724107, AA460041, AI215829, AI312833,
 de residue	AA772627, AA442303, AI936227, AI200468,
NO:2066, and where b is greater	AI282278, AI167870, AI130767, AW130869,
than or equal to a + 14.	AA43723
	AI338407, AI192747, AI283778, AI460353, W56676
	AA757574, N57307, AA676676, AI371859, AA992661
	, AI149595
	AA946707, AI245790, AI198433, AA831222,
	AI075992, AW073856, AI763210, AA442843, N21005
	AA526931,
	AA486260, AA024930, AA284849, N29407, AA768383
	1185523,
	, AI250412, AI269354, AA1331
	AA894509, AW170573, AA921691, AA284802,
	, AI862001,
	, AI186092,
	AA524571, AA229574, N95179, R62977, AA634150,
	g
	AI206465, AA640985, AW004616, AI016392,
	, AI262367,
	AI125021, AA143393, AI523228, AI339136,
	3, AI347544
	AI679670,
	AI033929,
	AI051474,
	AA635984,
	, N70831,
	AA296828, AA298518, R99897, AA298240, AI833094
	AA298536, AI186393, H22510, AI189398, H22509,

IR LOO 444 A L 44	HE8PBS6 B97898 Preferably excluded from the polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a net of a nucleotide sequence described by the general formula of a-b, where a nucleotide sequence described by the general formula of a-b, where a nucleotide sequence described by the general formula of a-b, where a nucleotide sequence described by the general formula of a-b, where a nucleotide sequence described by the general formula of a-b, where a nucleotide sequence described by the general formula of a-b, where a nucleotide sequence described by and nucleotide residues shown in SEQ ID NO:2067, and where both a and b nucleotide residues shown in SEQ ID A906505, A1948339, A167393, AA658034, A103336, A168037, and where b is greater and paragraphic, and where big seater and paragraphic, and seasons, and sea
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AAGGGD9, AW238652, AAG86218, AA482254, A4436333, AL291897, H99748, AT218514, AAGG0151, R67220, RA468385, AW302595, R82584, AI446687, AL283412, AL816752, H97740, AA53961, AU3394, R82285, AL68185, AW302595, R82584, AI446687, AM104955, AW261859, H21696, AI291596, H01942, AM104955, AW261859, H21696, AI291596, H01942, AAG18219, W93362, AA586370, AA28841, AA618219, W93362, AA586377, AW238247, AI190841, AA618219, W93362, AA58839, CIT106, AA297487, AA159156, A1472890, AA589549, AI492053, AA159555, AI472890, AA469417, AA076609, AA618219, AA531038, AA488520, AA29628, AA159555, AI41096, AI268122, AM239628, AA1365925, AI4109612, AA379712, AM379900, CIT487, AI36925, AI4910612, AA297272, AM379900, CIT487, AI36925, AA631038, AA188520, AA298674, AI36925, AA1910712, R2322, AA100383, AI572289, AA372564, R32172, AA583881, AA533195, R32216, R70940, A100172, R23223, AA100383, AI572289, AM372846, AA352092, AI890265, AA382912, AM372846, AA352092, AI890265, AA384494, W32530, AA288464, AA352092, AI890265, AA384494, W32530, AA215645, AI43731, AW266790, AI6930667, AI135545, AI43731, AW266590, AA65587, AA371033, AI135545, AI43731, AW266597, AA571033, AI135545, AI43731, AW266597, AA371033, AI135545, AI43731, AW266597, AR391033, AI135545, AI43731, AW266597, AI890265, AA394494, W32530, AI135545, AI43731, AW266597, AI890265, AA394494, W32530, AI135545, AI43731, AW266597, AI890265, AA394494, W32530, AI135545, AI43731, AW266597, AR3907610, AI1350112, AW384494, W32530, AI135545, AI43731, AW266597, AI990265, AA390687, AI135545, AI43731, AW266597, AB971033, AI155450, AI1359112, AI13401033, AI1417, AI15545, AI14731, AW266597, AA971033, AI155450, AI136414, AI1540667, AI107137, AI15545, AI14731, AW266597, AB971033, AI155450, AI144506, AI144506, AI100133, AI155450, AI144506, AI144506, AI100133, AI15545, AI14731, AI445069, AI007131, AI1033062, AI135945, AI144506, AI1350112, AW384444, W32530, AI1564500, AI14674, AI1570133, AI15545, AI147131, AA74507, AI1370133, AI1564500, AI149508, AI144506, AI1007033, AI1564500, AI144506, AI1007033, AI1007033, AI100703

AM191844, AL473451, A1922550, AL161279, AL748274, AM410102, A160610, Z97214, A2563, A12297, AF151109, 118978, X72644, E01573, E02319, 189947, AL050077, E06743, A23630, E12580, A08907, AR131821, AL17626, I17544, AL050155, AR068466, AL137480, U17594, AP028823, X66871, I33392, M27260, AL049233, A08913, I09499, AL137480, AL06911, A08911, AL050158, AR088484, AF031903, A08908, AF031147, AR10218, AR038854, AF031903, A08908, AF031147, AR10218, AR038854, AF031903, A08908, AR031147, AR03918, AR03912, AR08912, A08911, A08909, AL110218, AR03914, AL13775, AL117752, AR03913, L13297, I18355, S36676, E02253, AR03940, 149525, AR06875, S84466, U92068, AR183393, Y10655, AR117959, X76228, X87582, AR090344, AR117954, AL03576, AL13757, AR011880, U78525, AL110222, AL131648, X55446, AR107401, AL080148, AL137224, AL03548, AR068751, AL080448, AR0317, AR335, I19252, AR110269, AL137550, X98834, I08119, E15569, E02914, Y11254, AR0317, AR335, I19252, AR110269, AL03496, AR04373, S6735, AL131665, AL110269, AL09496, AR04373, S6735, AL131649, W09972, AL09496, AR04373, S6736, AL131761, AR090901, AR10224, I18979, S54990, A65965, AR117649, Y09972, AL090437, S784199, M88826, AL111749, AR090901, AR110269, AL080437, AR111849, M88826, AL111749, AR090901, AR110269, AL080438, AR111849, M88826, AL111749, AR090901, AR110269, AL108075, AR09090, AR090	91844, AI473451, AI922550, AI161279, 49274, AW410302, AI401697, AL023582, X70685 13019, A18777, AL080110, Z97214, A52563, 297, AF151109, I48978, X72624, E01573, 319, I89947, AL050277, E06743, A23630, 580, A08907, AF131821, AL117626, I17544, 50155, AR068466, AL137480, U77594, AF028823 871, I33392, M27260, AL049283, A08913, 499, AL137488, A58524, A58523, A12522, 18094, A08912, A08910, A08911, A08909, 10218, AR038854, AF031903, A08908, AF031147 39138, AF039137, A18788, S76508, AJ117394 50138, L13297, I18355, S36676, E02253, 392, U35846, S77771, I89931, AL117648, 080, AF039137, A18788, S76508, AJ137529, 29490, I49625, AR068753, S83456, U92068, 83393, Y10655, AF1177959, X76228, X87582, 15669, AL137294, AL137648, AS874, AF177401 80148, AL137294, AL096751, AJ005690, 37550, X98834, I08319, E15569, E02914, 254, A76313, A6136421, A5087943, 3750, X98834, I08319, E15569, E02914, 254, A76313, A6136421, A5087943, 3750, X98834, I03319, E15569, E02914, 254, A76313, AL137521, AF087943, 3750, AF184965, AL136842, X65873, AF08184, 3750, AR184965, AL136826, AL117649, X09972, 631, U73682, AL137521, AF090901, AF140224, 979, S54890, A65965, AB019565, A57389, 587, X84990, AF017152, D00174, AF112208, 80162, A65943, M92439, M80340, AC004200, 747, AF111112, M19658, AJ001838, A08456, 00937, AF118090, AF109155, AL110158,

736, 6623, 8, 008, 6, 6, 7,		Т93732,
ALI17457, ALI22045, ALI37284, ALI37533, U57352, X14634, I32738, A08916, E01614, E13364, A03736, ABD29065, AF069506, J05277, AF104032, ALI33623, S63521, ALI37478, U76419, ALI10221, AJ003118, AF185576, S79832, U42766, AF022363, I89944, D55641, I41145, X63410, ALI22110, AL049339, AF130470, ALI33640, AF013249, ALI37271, AF141289, AF017790, ALI33075, A07647, AF026008, X06146, AF00167, AL049382, AL080074, A70386, X61970, AF00167, AR055519, ALI37627, AF091084, AD337557, X79812, A77033, AT7035, X62580, AL049430, X95876, AL122100, AF043493, U87620, AF061795, AF090903, Y14314, AF151685, X99717, AF146568, AF090896	AF072128	AI818416, AI445972, AI923823, AA053602, AI798538, AI932810, AI521001, T93
i, AL137284, AL13753 18916, E01614, E1336 i, JO5277, AF104032, U76419, AL110221, A U42766, AF022363, I 3410, AL122110, AL0 i, AF013249, AL13727 AL049382, AL080074, AL049382, AL080074, AR055519, AL137283, AR017437, AL137283, A77033, A77035, X62 AL137461, E02349, A AL122100, AF043493, i, Y14314, AF151685,	AW385836,	AW237077, AW190814, AI1446688, AA135893, AI990405, AA346311, AI950541,
ALI17457, AL122045, AL137284, AL137533 Y14634, I32738, A08916, E01614, E13364 AB029065, AF069506, J05277, AF104032, S63521, AL137478, U76419, AL110221, AJAF185576, S79832, U42766, AF022363, I8 D55641, I41145, X63410, AL122110, AL04 AF130470, AL133640, AF013249, AL137271 AF141289, AF017790, AL133075, A07647, X61970, AF000167, AR055519, AL137627, A92311, AL133069, AF017437, AL137283, AL137557, X79812, A77033, A77035, X625AL049430, X95876, AL122100, AF043493, AF061795, AF090903, Y14314, AF151685, AF146568, AF090896	A1913916,	AA044768, AI826965, AI587426, AI587431, AI991706, AI803879, AI917076,
AL117457, AL12204 Y14634, I32738, A AB029065, AF06950 S63521, AL137478, AF185576, S79832, D55641, I41145, X AF130470, AL13364 AF1289, AF01779 X06146, AF000167, X61970, AF000167, A92311, AL133069, AL137557, X79812, AL049430, X95876, I17767, AL137554, AF061795, AF09089 AF146568, AF09089	AA345449,	AA044731, AI989722, AA053091, AA112375, AI493214, AI984082, AI582971, AI611349,
	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 251 of SEQ ID NO:2068, b is an integer of 15 to 265, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:2068, and where b is greater than or equal to a + 14.	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 760 of SEQ ID NO:2069, b is an integer of 15 to 774, where both a and b
	898087	898136
	HTPGE66	HWLIL19
	2068	2069

			correspond to the positions of	AI434008, AI913316, AI932552, AI431343,
			nucleotide residues shown in SEO ID	
			and where b	
			than or equal to a + 14.	
2070	HPJEE80	898157	Preferably excluded from the	AA314262, AI698145, AI751509, AI765378,
			present invention are one or more	AI819921, AI309793; AI983094, AI889488,
			polynucleotides comprising a	AI691017, AI478725, AI418367, AI768787,
			nucleotide sequence described by	AI336867, AA770272, AI579948, AI347373,
			the general formula of a-b, where a	AA773349, AA287318, AA187540, AA854659,
			is any integer between 1 to 2606 of	AI637840, AI566584, AA305439, AA451739,
			SEQ ID NO:2070, b is an integer of	AA287399, AA255886, AA689402, AI961717,
				AI624071, AW444697, H24906, R59469, AI636153,
			correspond to the positions of	AL037168, AW151230, AA256684, AA694475,
			nucleotide residues shown in SEQ ID	AI861989, H02063, H26485, H13596, AA256683,
			NO:2070, and where b is greater	AA348853, AA336954, H02078, H44525, AA354340,
			w.	Z43173, AA337732, AI565023, H44530, AW297887,
		_		R75751, H26324, AA336921, R41517, AA775352,
				σ
				AI954448, AA336373, AA336703, C02323, AW391166
				AI858347, AW379208, AA634601, AA449368,
				AI611218, AA262646, AI860650, AA282616,
				AL119399, AL119457, AL134524, AL119324,
				AL042544, AL119443, AW392670, AW372827,
				AL119391, AL119464, U46346, AL134902, AW384394
				AL042614, AL119319, AW363220, AL119484,
				AL119497, AL119335, U46341, U46350, AL119341,
				U46347, U46351, AL119439, AL119444, AL119396,
			-	AL119483, AL119418, AL119496, U46345, AL134518
				AL134528, AL037205, AL134525, AI142132,
				AI142137, AL134538, AL042970, AL042450,
				AL042965, AL042975, AL134529, AL042542,
				AL043019, AL042984, AL043029, AL042551,
				2319, 29
				, AR060045, AL035687, Z65447, AB02643
				AR060234, AR066494, A81671, AR054110, AR069079

				AR043113
2071	HWLOX67	898192	Preferably excluded from the	AL120532, AI587307, AI093091, AI769686,
			present invention are one or more	AI050667, AI372945, AA250932, W15253, N49198,
			polynucleotides comprising a	W39173, AA894448, AA975408, Z21307, AA846588,
			nucleotide sequence described by	AC002554, Z73358
			the general formula of a-b, where a	
			is any integer between 1 to 1462 of	
			SEQ ID NO:2071, b is an integer of	
			15 to 1476, where both a and b	
			correspond to the positions of	
		_	nucleotide residues shown in SEQ ID	
			NO:2071, and where b is greater	
			O)	
2072	HCRNK75	898355	Preferably excluded from the	, AI823427, AI377127
			present invention are one or more	AW168810, AA293513, AW088676, C17686, AI289654,
			polynucleotides comprising a	AI207850, AI890720, AI805626, AI824271,
			nucleotide sequence described by	AI344359, AI300131, AA574103, AI686750,
			the general formula of a-b, where a	AA315866, AI709243, AA252863, AA585439,
			is any integer between 1 to 2210 of	AI758734, AW375857, AA348962, AI525556,
			SEQ ID NO:2072, b is an integer of	AA585453, Z28355, AA585440, AI525316, AI535639,
			15 to 2224, where both a and b	AI541510, AI546855, AA336552, AI541374,
			correspond to the positions of	AI556967, AI525328, AI541514, C15189, AI541523,
			nucleotide residues shown in SEQ ID	Z30131, AI526180, AI546999, AI541534, AI525306,
			NO:2072, and where b is greater	
			than or equal to a + 14.	, AI541365, AI382291,
		-		AI541017, AI525431, AA585356, AI557731,
				_
				AI541317, AI546945, AI535813, AI557799,
			-	AI540967, AI557262, AI525653, AI541508,
				AI541307, AI541535, AI557082, T11028, AI546899,
				D61254, R29445, AI557787, R28735, AI546875,
				AI541205, AL040510, AL040625, AL045817,
				AL041238,
				AL040322, AL041131, AL046330, AL041051,
				AL041292, AL040119, AL047036, AL047170,
				AL047057, AL047219, AL041227, AL040463,

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AL041197,	AL041096,	AL041163,	AL041324,	AL043496,	AL041159,	AL036500,	AL040252,	AL040091, AL040128, AL040168,	AL040342,	AL045684,	AL043677,	AL040149,	AL041602,	AL040253,	AL040458,	AI525320,	AL040329,	AL040148,	AL046392,	AL043537,	AL039316,	AL043848,	AL044258,	AL040768,	AL046994,	AL046914,	AL049007,	AL043468,	AL039744,	AL044015,	AL037341,	AL045991,	AL045671,	AL041168,	AL041246,
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		07, I13349, A91965, I66498, I664
	-	66494, I66487, I66497, I66496,
	-	66486, A83643, I66485, I66488,
		66491, I66492, I66493, A83151,
		84, X81969, A25909, AR062871
	-	I18895, A85395, A85476, AR062872, AR062873,
		, AJ244005, AJ244003,
	-	86,
		A43188, A91752, I63120, A98767, A93963, A93964,
	_	, A98423, A98432, A98436,
		, Y16359, AR038762, I4468:
	_	A86792, X83865, A84772, A84776, A84773, A84775,
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		, I15718, A58523, E03627,
		I8455
		, A60111, A23633, AR007512
		, IO5488, I61310, A60209,
		, A60961, A60977,
		, A11178, E01007, A10361,
		, AR027318, A68112
		, A23997,
-		, A08030, A20502, I62368, A35
		v,
		9, AR043601
		, A60990, A47368, A60987, Il9
		.11, I6
		52, AR009151, I63561, I63563,
		A02710, AR035193, E14304, A
	_	
		27396, AR027100, I44531
		44516, A70040, E16678, AE
		I08776, I15353,
		AR068510, AR068509, A63954, I91969, I26929,

				TAMETE 136030 136030 136037 158333 158333
				, 125041, A24783, A24782, A92133
				A95117, A90655, A38214, I56772, I95540, A95096,
				A95106, A95105, AF149828, I01995, I08051,
				AR031566, I60241, I60242, AR038066, A20699,
				E00696, E00697, E03813, AR027099, Y09813,
				AR051652, AR051651, Z32836, AJ230935, D50010,
				AJ230902, AR035975, AR035974, AR035977,
				I05558, AJ230972,
				A91754, AR031374, AR031375, AR020969, A92666,
				A92668, A92667, A92665, E12584, AJ230951,
				A70872, AJ231009, A22738, I08389
2073	HOGDR01	898418	Preferably excluded from the	AI940071, AW383315, AW383305, AW383297,
			present invention are one or more	AL134527, AW384394, AW363220
			polynucleotides comprising a	AL119443, U46347, AL119522, AW372827, Z99396,
			nucleotide sequence described by	AL119319, AL119324, AL119457, U46350, AL119439,
			the general formula of a-b, where a	U46349, AL119484, AL119391, AL043003, AL119483,
			is any integer between 1 to 806 of	AL119497, AL119401, AL119363, AL119444,
			SEQ ID NO:2073, b is an integer of	AL119355, AL119396, AL134525, AL037205, U46341,
			15 to 820, where both a and b	AL134531, AL134902, AL042984, U46346, AL119418,
			correspond to the positions of	AL119399, AL119335, AL042542, AL134538,
			nucleotide residues shown in SEQ ID	AL043019, AL042544, AL042965, AL042975, U46345,
			NO:2073, and where b is greater	AL042614, AL043029, AL042989, AL042450,
			than or equal to a + 14.	AL042551, AL119464, AC003965, AB026436,
				AR069079, AR066494, AR060234, A81671, AR054110,
				AR043113
2074	HHATR06	898427	Preferably excluded from the	AI478733,
			present invention are one or more	AI379565,
			polynucleotides comprising a	AW294114, AA427646, AI751750, AA594137,
			nucleotide sequence described by	AA947297, W95460, AI057073, AA405402, AA788855,
			the general formula of a-b, where a	AW068453, AW068711, AW177719, AI341112, H73236,
			is any integer between 1 to 1473 of	AW167569, AA232452, AA427487, AA041328, W95567,
			SEQ ID NO:2074, b is an integer of	AI652166, AA853047, H74164, R34003, AA041304,
				W02069, AI341381, AW192052, AA580289, AL119457,
			correspond to the positions of	AL042544, D30965, D31176, AL119324, AL119399,
			nucleotide residues shown in SEQ ID	AI918637, AL046052, AL042866, AI690472,

_	NO:2074, and where b is greater	AI918408 AL045891 AI689380 AI433206
	00000	COLK LACALCTK COLLCOME COC
	⊣ + ਚ	, AWOZ4/33, AI343281, AIU96 , AMOZ4/33, AI343281, AIU96
		, AIZ41884, AI3/1228, AIS8291
		, AI446405, AI564160, AI91
		AI273919, AA838230, AW083489, AI865942,
		AW194441, F36003, AI499104, AI887775, AW151974,
		, AW058275,
		84, AI634930,
		AW008253, AI686081, AI921922, AA749024,
		AI125845, AI472476, AW085866, AA480074,
		AI313320, AW022494, AI313352, AI310920,
		AI307503, AI671284, AW020288, AI612732,
		AI933926, AI336585, AI334913, AI349266,
		_
	-	AI312146, AI312339, AI309431, AI340537,
		AI312165, AI345258, AI349288, AI349628,
		AW196105, AA835966, AI340610, AI307459,
		AI343140, AI349971, AW168693, AI307507,
		AI348879, N22406, AI340639, AI311604, AR035969,
		L24896,
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		M30514, AF
		2, M79462, AL133629,
		AL137658, AL110280, AR011880,
		AL049464,
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		AL137665, A90832,
		F161699,
		AL096720, Y11435, AF113694, X54971, Y10080,
		3, AF051325
	-	
		AL110171, Y10655, AF118064, AL049314, AL137558,
		L31396, U68387, AL137656, AF010191, L31397,
		AF151109, AF140224, AL110159, X76228, S63521,
		U92068, AF148129, AF081366, Z72491, S69385.

				157350 Y92070 BE026124 H57352 V14634
				AL137267, Z48796, AC007458, AF017
				AL133636, S61953, L78810, AF213396, U67328,
				AF114818, AF113676, AL137534, AF016271,
				, S75997, A
				S73498, AF118558, E04257, AR005011, U80919,
				AP000130, AP000208, AP000247, AL035458,
				AF144700,
				AL080158
2075	HLQDM07	898541	Preferably excluded from the	AI806250, AA455382, AI084580, AW368035,
	,		present invention are one or more	AA005065, AI088155, AI566044, W92235, AA706063,
			polynucleotides comprising a	W92236, AA299662, AA004847, H56718, T77776,
			nucleotide sequence described by	AA002009, AA227236, AI922495, AA722941,
			the general formula of a-b, where a	AA456022, AA299663, AA001788, H56641, AL119457,
			is any integer between 1 to 2372 of	AW392670, Z99396, AL119319, AL119355, AL119324,
			SEO ID NO:2075, b is an integer of	AL119497, U46350, U46351, AL119363, U46349,
				AW372827
			correspond to the positions of	AL119341, AW363220, U46347, AL119484, AL119443,
			nucleotide residues shown in SEQ ID	U46341, AL119444, U46346, AL119439, AL119522,
			NO:2075, and where b is greater	AI142134, AL119396, AL119335, AL043033,
			equal to a + 14.	AL037205, AL119401, AL134538, AL134542,
				AL134528, AL134902, AL134531, AL134533,
				AL119418, AL119399, AL042984, AL119496,
				AI142132, AL134525, AL134536, U46345, AL119464,
				AL042450, AL042614, AL043029, AL042544,
				AL043011, AL043019, AL042542, AL042965,
				AL042975, AL043003, AL042551, AL132826,
				AF169677, U42975, AB026436, AR066494, AR060234,
				AR054110, A81671, AR069079
2076	HDPBW68	898651	Preferably excluded from the	AI797914, AA232727, AI264354, AA242826,
			present invention are one or more	AI373844, AI421152, AI693559, AA293798,
			polynucleotides comprising a	AA242961, AI681069, AA987481, AA253496,
			nucleotide sequence described by	AA865918, AA394280, AA699441, AW193319,
			the general formula of a-b, where a	AA534330, AI246675, AI690035, AI921391,
			teger between	AI696791, AI696792, AI962498, AA478182,
			SEQ ID NO:2076, b is an integer of	AA845215, R02588, AA501984, AA253392, AA975909,

			15 to 3893, where both a and b	AI359321, R02707, AI
			correspond to the positions of	AA065210,
		_	nucleotide residues shown in SEQ ID	AI217878, AI470976, AI640699, AL119324,
			NO:2076, and where b is greater	AL119457, AL119399, AL042544, AL119443,
			equal to a + 14.	AW392670, U46346, AL119355, Z99396, AL134525,
		_		U46351, AL119319, U46349, AW372827, AL119483,
				Ċ,
				U46350, U4
				AL119444, AL119341, AL119418, AL134902,
				AL119439, AL119335, AL119522, AL037205,
				AL119396, AL119401, AL134538, AL134527,
				AL119464, AL042450, AL043033, AL042984,
				AL119496, AL134536, U46345, AL042433, AL042614,
				AL043029, AL043011, AL043019, AL134542,
				AL042542, AL042965, AL042975, AL043003,
				AL042551, AF113925, AF126484, AF149774,
				AC006027, AB026436, AR060234, AR054110,
				AR066494, A81671, AR069079
2077	HISCI15	898814	Preferably excluded from the	L44393, AA434356, AI524406, AW062354, T31737,
			present invention are one or more	H14980, Z43676, N40577, R08471, N25869,
			polynucleotides comprising a	AA256007, N41934, N28530, AA808513, T92387,
			nucleotide sequence described by	R02302, AW383005, AB011165, AF117754, AR022169
			the general formula of a-b, where a	
			is any integer between 1 to 3219 of	
			SEQ ID NO:2077, b is an integer of	
			15 to 3233, where both a and b	
			correspond to the positions of	
			nucleotide residues shown in SEQ ID	
			NO:2077, and where b is greater	
			than or equal to a + 14.	
2078	HCYBH77	898946	Preferably excluded from the	7, AW268365, AI433801,
			present invention are one or more	AW376970,
			polynucleotides comprising a	AI744244, AA179345, AW264850, AW239439,
			nucleotide sequence described by	
			the general formula of a-b, where a	AI270669, C18854, AA186804, AA505958, W63641,
			is any integer between 1 to 2967 of	W52261, AL036582, R50884, H17527, AA033538,

			SEO ID NO:2078, b is an integer of	AL048651, AW149146, AA305384, AW273640, R50765,
			, where both	C17088, AA356773, AI698410, R07093, AA134840,
			correspond to the positions of	AI985957, AA808140, AA367305, W79703, AA381398,
			residue	AF123887, AF144695, AR018794, AR018857
0000	170 11011	000	udi to a + 14.	TEOCOLIA COCESOAN ETITOOMA CICOCEA
6/07	HPJAS61	051668	excidaed from the	, AWOU/113, AAO36262, A13020///
			()), A14169/8, Aw2/5894, Aw23694.
			tides comp	AW167603, AI031828, AI624036,
			nucleotide seguence described by	N63417, AI
			the general formula of a-b, where a	AI075944, AI347803, AL134813, AA010795,
			is any integer between 1 to 2444 of	AI991823, AA608692, AW188444, AI765847,
			SEQ ID NO:2079, b is an integer of	AI580486, AA488368, N38923, N30935, AI093100,
			15 to 2458, where both a and b	AI453400, AI434592, AI300853, AA457119,
			correspond to the positions of	AA455498, AI880713, AW050861, AI274340,
			nucleotide residues shown in SEQ ID	AI309910, AW207240, AA633538, AI188595, H98907,
			NO:2079, and where b is greater	AI308095, AI863003, AA705931, AA165111,
			equal to a + 14.	AI066618, AI261549, AI470214, AI282600,
				AI635033, AA011134, AA583904, N95694, AA973598,
	_			
				AA599424, AI864628, AI831364, AI610395,
				AI245485, AA649888, AI672081, N72372, AA293614,
_				N95723, H77346, AI270457, R53634, AA829048,
				AA062785, AA479044, AA826668, T65751, H58487,
				H81750, AI092643, AA190410, AW300733, AW264761,
				w
				~
				AI925804,
				AI309181, AA627576, AA430543, AA430544, R87874,
				AA369400, H77345, AA468680, AA853269, N52644,
				AA130245, AA157200, AI160148, AA834736,
				AA705668, AI124918, AA156892, AA948320,
				AI609381, R45075, AI701123, AW178256, AA376537,
				AA296785, AA190800, H52032, AI673683, H57644,

A	AA477526, AI401060, N35904,	3, 1916819, 132, 13, 1, 1,		, , A1479577, AL121496,	, A74674, 3249, 92, 5604, 7620, AL133049, AF119336, AF118090,
	~ ~	A297348, 246, AA916 5, W70132, 1590043, 1559752, 1621341, 79232, AI6	1680467, M089844, L042365, 1491852, W327527,		944 8468 333 077 107 8,
MA91349, T55826, AND A455499, AL623320, AND A5359, AL623260, AND A163667, AL624515, AL63667, AL64544, CO1875, AL64518, A	AA AI ,	.IS87427, p 90317, H85 911, N4649 W075382, p .I333104, p .I284517, p		1445611, A 1475371, A W128834, A 1683395, A 1890223, H 66465, A12	42, AF200192, A 71, AF051150, A 189947, X9349 A77033, A7703 S36676, AL137 AL080140, AR0 AF111849, AB0 AF11849, AB0 AF158248. T
N94355, A44 A4636499, A4654699, A A163667, A A163667, A A163667, A A163667, A A163667, A A163860, A A163860, A A163860, A A163131, A A169108, A A169108, A A169108, A A169109, A A169109, A A169391, A A16369415, A A16369415, A A16369415, A A163694, A A163694, A A163694, A A163694, A A163694, A A163695, A A163699, AB	5 64	σ σ		A514684, A L047344, A I341838, A I926330, A I884303, A 75779, AI8 WI18553, A	FC00342, A FC050171, A 49466, I89 29065, A77 70386, S36 66871, AFO 17587, ALO 17437, AFI L133088, A
	→	&			AF117892, AF200342, AF117892, AF050171, 148978, AL049466, I8 109499, AB029065, A7 AB026995, A70386, S3 AR029580, X66871, AF A91162, AL117587, AI U49908, AF017437, AF AF111851, AL133088, AF022813, AL137558.
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	U72621, U89295, AR038854, AL050208, AL133062,
	AL133010, AL137480, Z97214, AL050092, AF079763,
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	U35846, AF102578, AL110
	AL137271, AF061795, AF151685, AL122121,
	vo
	i, U75932, AL117648, Y10655, AF1130
	L13746
	AL137530, AF200464, E06743,
	AJ000937, AL137560, X80340, AF141289, AF185614,
	AL117435, U51587, AL137627, X76228, AL137557,
	AL117416, AF026124, AL1221
	AB029066, L13297, A08907, AF111112, AR020905,
	J05277, A41575, A65340, Y10936, E12580, U83980,
	A08912, AL110221, AF090900, U73682, AR068466,
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	AL122050, AL137258, AL080234
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	AF131821, X89102, AF183393,
	I68732, A08911, A18788, I89931, X97332, L04849,
	_
	89934, IO83
	AF087943,
	AF068615,
	, AL137284
	A58523,
	U53505, L30117, AF047716,
-	75, U62966, AL117440, AF185576, AF047
	L04504, AL137463, AF182215, S61953, AL137657

HCRMK25 899224 Preferably excluded from t	Preferab	آ و	from the	AA704087,	AW373819,	AW380680,	AI752796,
	60	60	ng a	AA428419,	AA780507,	A1906013, AA668306,	AW385383, AL036004.
nucleotide sequence described by	nucleotide sequence descr	nucleotide sequence descr		AA600085,	AI751526,	AI751512,	- rV
the general tormula of a-b, where	the general tormula of a-b	the general tormula of a-b	, where a	AW239513,	W49750, A	W49750, AA773949, N36271,	36271, W63574,
SEO IN MO: 2000 P in interest	GEO ID NO: 2000 P in the	SEO ID NO: 2000 h in the	2636 of	AA780819,	AA457563,	AI753606,	AA464937,
.coo, n is an i			eger or	AA454895,	AW385419,	AI905876,	AI752292,
13 LO 2030, Where both a an	13 LO 2030, Where both a an	13 LO 2030, where both a an	and b	AA181456,	AW068389,	AI751743,	AA457359,
correspond to the positions of	collespond to the positions	correspond to the positions	í	AI751229,	AI752349,	AI365966,	AA293647,
re residues show	ນ	ນ	n SEQ ID	AA554805,	AI752176,	AI751283,	AA489941,
, and where b	and where b is	and where b is	ter	AA457511,	AI751586,	AA788961,	AW352231,
than or equal to a + 14.	ednal to a +	ednal to a +	-	AI752829,	AA487731,	AA789233,	AI750701,
				AI752337,	AA487393,	AW373901,	AA457430,
				AA704140,	AA457469,	AI905974,	AA169848,
				AA703999,	D79055, AI	752205, AA	AI752205, AA434290, AA489933
				AI752293,		AA434353,	AA489957,
				AA780675,	AW352222,	U53087, AI	
				AI752797,	AI752212,	AI751798,	AW373788,
	· ·			AI752270,	AW373787,	AI925580,	AI752737,
-				AW373833,	AA121851,	AA456983,	AI752171, N34179,
				AA458778,	AA454883,	AI751523,	AA679516,
						AW393626,	AI751886,
				~	AW384994,	AI751927,	W24625, W00702,
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				AI751476,	W52302, R7	W52302, R71009, AW373902,	3902, AA486177,
-					AA961963,	AA594126,	AA476858,
			_	_			AI751810,
				ω		8	AI750740,
				ς, ,		R73275, AW	AW068267, AW373874,
			_	_	AW373808,	8	AI750278,
				AW373834,	AA136731,	0	AI906084,
					AA359001,	. ~	AA780557,
				AA453844,	AA318038,		AA668143,
					AI751652,	,	AW366380,
				62,	W24650, AA477811,		AI963017, AA293756
				H53916, AA	AA169864, AI		

	AW362721, AA373886, H82181, AA434473, AA334411,
	AI922681, AI963366, AI752830, AA457291,
	AA668375, AA443350, R84909, AW067859, T29584,
_	AW385969, AA339992, AA379018, AA326804,
	AA378055, AA375369, AI752739
	, AA376383, AA359377
	AW363460,
	W00543, AI750767, AI
	AA373229, AA218722, AA853386, AA332082,
	AA359277, AW384999, AA372196, AA333869,
	AI796681, AL035880, AW082115, R69349, AA359183,
	AA359695, AA852609, AA366521, AA434079, T49549,
	AA853491, AA669422, AA377936, T53285, AA377860,
	AA595560, AA346953, AW068393, AA852626,
	, AA256215,
	AA070541, T4
	, AA339830
	AR048312, AB015438, AB008373, U03419, M14423,
	X98705, S67482, M17491, X06269, AF169346,
	9, Y15918, D83228, Y15919
	X98708, Y15913, Y16346, J00112,
	, Y08643, Y15916, J00111,
	, T53375, T99669, R01522,
	H50793, H52341,
	W05288, W05816, W25354, AA167235, AA167584,
	AA987726, AA
	, AA853652, AA853657, AA8
	AA853790, AA852117, AA852484, AA852780,

			AA852811, T49210, T49936, D45437
HNTRVII	1 899632	Preferably excluded from the	AI192806, AI636301, AW070460, AI264134,
		present invention are one or more	AI808610, AL047490, AW337234, AW272771,
		polynucleotides comprising a	AA621722, AA902441, AW338001, AI572907,
	-	nucleotide sequence described by	AW088299, AA630592, AW241806, AW338392,
		the general formula of a-b, where a	AW119186, AW361987, AI598101, AW079856,
		is any integer between 1 to 2288 of	AI932992, AA314261, AI380908, AI571554,
		SEQ ID NO:2081, b is an integer of	AA431144, AW362042, AI741945, AW029103,
		correspond to the positions of	AI246132, AA188213, AI092692, AI129947,
			AA969200, AA495870, AA774660, AA835498,
	·	NO:2081, and where b is greater	AA825370, AA432163, AI520696, AI624063,
		than or equal to a + 14.	AI026883, AA888774, AA186360, AW390429,
			AI692914, AA262302, AA156547, AI289833,
			AI678753, N76487, AA676856, AA190635, N36869,
			AA512918, AI392858, AI571545, AA262303,
			_
	_		2, AI129465,
			H48412, N94959, AI218172, AI221051, AA577253,
			AA086067, AI439435, AA112358, AI241626, R80350,
			Q.
			AI537627
			AA192529, R77146, AA188562, AI250628, H73378,
			, H45701, AI281554,
			, H56566, AI445365
		•	4, W19537, T78819, H45752,
			47345, AA973983, R62945,
_			T60098, R45931, H98238,
			7, R70102, H12066, R354;
			H03315, AA369106, W25341, R80240, AI263665,
			AI803872, AA757310, AI
			R76607, R67545, AA622166, H16044, T82361,
			AI802973, AA188660, F077
			H54185, H03316, F08108, R62997, T94841,
			AW338108, T94886, AL045149, H97241, AA630804,

				AA344563, F02937, AW316643, AI635890, H71561, R70103, AI985724, N27010, AA2 N72946, R76608, AW366579, N49618, T73 AI587589, M31516, I41330, I05091, I09 M15799, U88576, S67775, M30142, I0921 A65264, AR031710, AR066586, AR066589, M64356, S51407, AB003312, AB003313, A AB003316, AR016512, AR016513, Z63791, I64711, A AR016516, I64714, M64652, AB003319, A C7265	, F02937, AW3166- R70103, A1985724 R76608, AW366579 , M31516, I41330 U88576, S67775, AR031710, AR0665 S51407, AB003312 , AB003317, AR01 , AR016513, Z637	F02937, AW316643, AI635890, H 0103, AI985724, N27010, AA218 6608, AW366579, N49618, T7366 M31516, I41330, I05091, I0921 8576, S67775, M30142, I09216, 031710, AR066586, AR066589, A 1407, AB003312, AB003313, AB0 AB003317, AR016514, AB003315, AR016513, Z63791, I64711, AR0 I64714, M64652, AB003319, AB0	3, F02937, AW316643, AI635890, H56567, R70103, AI985724, N27010, AA218591, R76608, AW366579, N49618, T73663, 9, M31516, I41330, I05091, I09215, U88576, S67775, M30142, I09216, I05094, AR031710, AR066586, AR066589, AF052110, S51407, AB003312, AB003313, AB003314, 6, AB003317, AR016514, AB003315, 2, AR016513, Z63791, I64711, AR016518, 6, I64714, M64652, AB003319, AB003318,
2082	нwLоизз	899644	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 1944 of SEQ ID NO:2082, b is an integer of 15 to 1958, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:2082, and where b is greater than or equal to a + 14.	AL037051, AL045353, AL039386, AL036973, AL038837, AL037615, AL036238, AL03765, AL037027, AL037027, AL037027, AL037027, AL037027, AL037027, AL047064, AL044064, AL044064, AL044064, AL044159, AL041133, AL041133, AL041133,	ALO40992, ALO39423, ALO38531, ALO39659, ALO39659, ALO37639, ALO3767, ALO3735, ALO3735, ALO3735, ALO3735, ALO40052, ALO40052, ALO40052, ALO40052, ALO40052, ALO40052, ALO40052, ALO40052, ALO400052, ALO400052, ALO400052, ALO400052, ALO400052, ALO400052, ALO400052, ALO400052, ALO400052, ALO4000000000000000000000000000000000000	AL042909, AL039128, AL03726, AL03726, AL039674, AL039678, AL039678, AL03727, AL049018, AL049018, AL049018, AL049018, AL04186, AL04186, AL04186, AL041635, AL041635, AL041635, AL041635, AL041635, AL041635, AL041635, AL041635,	AL039109, AL045337, AL038025, AW235098, AL039629, AL039629, AL039566, AL039566, AL039566, AL040576, AL040576, AL041314, AL0413814, AL0413814, AL044037, AL044037, AL044037, AL044037, AL044037, AL046625, AL044074,
				AL041602, AL043441,	AL045472, AL045671,	AL046442,	AL036158,

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	VETECOTIVE POINT	1 0		,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,	
-	AD045467,	ALU44236,	ALO40444,	ALU442/2,	
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	AL044274,	AL040745,	AL040463,	AL047183,	
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	AL040571,	AL046327,	AL044165,	AL040091,	
	AL045817,	AL041131,	AL040090,	AL047012,	
-	AL047057,	AL041292,	AL041051,	AL040168,	_
-	AL041346,	AL037341,	AL041955,	AL040414,	
	AL043775,	AL041096,	AL039744,	AL046330,	
	AL041197,	AL045989,	AL047036,	AL040553,	
	AL040253,	AL040155,	AL040082,	AL039360,	
	AL045857,	AL036117,	AL040329,	AL041358,	
	AL043538,	AL041163,	AL041324,	AL036725,	
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	AL043941,	AI906064,	AL041278,	AL040255,	
	AL038043,	AI634028,	AI906040,	AL040621,	
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	AL045725,	AL039915,	AL041140,	AL043612,	
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	023	AL037085,	AL038821,	AL046147,	
	AL038761,	AL041233,	AL036679,	AL134524,	
	AL036152,	AL041246,		T24119, AL03908	5,
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	AL039416,	H00069, AI	AL041347, AJ	AL036733, AL036900	,

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I40851, A60983, I60241, I60242, A02710, E12615,
I28266, I21869, AR036903, A70040, I66498,
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157, AR054109, A8
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AJ244004, AR022240, A85476, AR038762, A85395,
X68127, AR031374, A49700, AR031375, A58521,
AR020969, AR025207, AR036905, A38214, A44171,
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S, E16678, E16636, I44516, A82653, A9301
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AS8525, I49890, AF1562 AS8526, A91753, M28262 , I18302, AJ244005, AJ , A67220, A90655, Y119 100074, D88984, I03665 , I13521, I03664, I520 AJ244007, I66485, I489 AR038286, I25041, I924 , AR008430, I19525, E0 , A68112, A68104, A602 A60211, I15717, I15718 A77095, I08396, I07429	AIO81543, AW024140, AA742572, AW327486, AA593322, AI239527, AI362956, AA977531, AA865071, W76539, AA988767, AI240922, W56688, acribed by AW406326, F25349, W56696, AIS90417, AA773777, a-b, where a N80724, AW273295, N72158, AA356111, AA588352, I to 1233 of F36934, T23069, AA594466, AI002202, AW410884, I integer of AA779395, D80166, C14331, C14429, D80038, and b AA779395, D80166, C14331, C14429, D80038, bwn in SEQ ID D80227, D80195, D51799, D80269, D58283, D59859, D51423, D5927, D80195, D51799, D80193, D80196, D80188, D59927, D80219, D59502, D81030, D59889, D57483, D80022, D80366, D59610, D80378, D80045, D50979, D80164, D50995, D80241, D59787, D80024, T03269, C75259, C14014, D59467, C15076, C13389, D51026, D80168, C14227, W21835, D81111, D51079, AA305578, AIJ8893, D80111, D51022, AW179328, AA51188, AW178775, AW378532, D80522, D59695, AA51188, AW178775, AW377671, AI905856,
	preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where is any integer between 1 to 1233 SEQ ID NO:2083, b is an integer of correspond to the positions of nucleotide residues shown in SEQ NO:2083, and where b is greater than or equal to a + 14.
	899661
	HAPNO50
	2083

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	AW352117, C05695, AW176467, AW375405, AW378540,
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	AW179024, D59373, D80247, AW177505, D80014,
	AW179020,
	AW177733,
	AW178754,
	AW352174, AW179004, AW179012, AW178914,
	AW177722, AW177728, AW367967, AW179009,
	AW178774, AW178911, AW378543, AW352163, C06015,
	C14344, AI535686, AW178983, AW352120, D80258,
	AW378539, D59627,
	3
	D59317, H67866, AI535961,
	A84916, A62298, A62300, AJ132110, Y17188,
	AR018138, X67155, A67220, D89785, A78862,
	46
	I82448, AR016808, AB012117, A30438, Y12724,
	A86792,
	œ
•	A45456, AR066488,
	AR052274,
	R038669, AR066487, AR066490
-	, U46128, D88507, AR064240,
	AR016690, I18367, D50010, AB033111, A63261,

				AR008408, AR062872, A70867, I79511, D13509,
				A64136, A68321, AR060133, U87247, AB023656,
2084	HBSAK60	899776	Preferably excluded from the	R28735, R29445, R45895, AA585325.
			present invention are one or more	3, R29657,
			polynucleotides comprising a	AA585101, AA283326
			nucleotide sequence described by	D57491,
			the general formula of a-b, where a	AI546971, AA585439, Z32822, Z2
			is any integer between 1 to 2115 of	
			SEQ ID NO:2084, b is an integer of	
			15 to 2129, where both a and b	AI541365, AI541013, AI525500, AI557740, C16305,
			correspond to the positions of	ø
			nucleotide residues shown in SEQ ID	AI547250, D59751, C15406, D54897, D53161,
			NO:2084, and where b is greater	AI546945, AI541374, AI525306, AI525856, D53447,
			than or equal to a + 14.	AI541205, AA585155, C16292, AI526078, AI541517,
				AI546996, D55233, AI557731, AI525431, C15069,
				AI541535, AI547039, AI526184, AI525556, Z32887,
				AI525316, C16294, C15120, Z30131, D52835,
	-			AIS57727, R29177, AI526194, C15737, R29179,
				C15762, AIS41346, AIS57807, AIS46891, AIS41523,
				AIS57084, D57186,
				, AI547202, AI526191
				AIS57408, R29172, AIS57155, D60730, AIS57602,
				н
				, AI557808
				AI535660, T41289, AI526180, AI546829, AI535639,
				AI526109, AA174170, AI557039, AI540903,
				AI526195, AI547137, AI541422, T41329, Z33559,
				AI524904, AA514191, AI526024, AI526158,
				AI525656, AI526112, AI557533, AI525286,
				, AI541510, AI541345, D51433,
				, AI546831, AI525332
				AI541027, AI557264, D59458, AI541415, C14723,

				AI557238,	AIS57852, C14322, C1	C14391, AI557799,
				AA585434,	AI526205, AI540882,	AIS41390,
				AI541017,	AI524890, AI547189,	AA585117,
	•			AI526117,	AI546954, AI541353,	AIS41508,
				AI546901,	AI526187, AI557082,	AA585430,
				AI557285,	AI557041, AI541492,	AI524891,
				AI547026,	AI557796, AI541515,	AI557786,
				AI557317,	AI525076, AI525114,	AI525168,
	70 <u>.</u>			AI540944,	D61060, AI557810, C1	C14210, T10982,
				AIS47071,	AI541410, AI541423,	AI541075,
				AI525653,	AA585420, AI557802,	AIS57785,
				AI046024,	AI526169, AI526144,	AR038855,
	•			AR062871,	A25909, Y09813, Z328	Y09813, Z32836, AR054723,
				AJ244005,		0010, D13509,
	-			AJ244004,		3872, AR062873,
	•			A20700, D	A20700, D78345, A43189, A43188, AR017907,	3, AR017907,
				AR038762,	AJ244003, A98420, A98423,	8423, A98432,
				A98436, A	A98417, A98427, X82786, X55486, X76012	;, X55486, X76012,
				AC005913,	A98767, A93963, A93964, I63120,	64, I63120,
				AJ244006,	AJ243486, AR031365, AR003381,	AR003381,
				AR031358,		
2085	HDPOD73	998668	Preferably excluded from the	AA478514,	, C00579,	AI708851, AI581139,
			present invention are one or more	AA640563,	R81679, AA367920, AI	AL046227, AI433131,
	•		polynucleotides comprising a	AI754257,	AW117882, AI242236,	AF113694,
			nucleotide sequence described by	AC004813,	AP000347, AL035587,	Z95114, AC004883,
			the general formula of a-b, where a	AC005291,	AF091512, AC004383,	Z82206, AP000344,
		_	is any integer between 1 to 774 of	AC004987,	AC006013, AF090900,	AC005274,
			SEQ ID NO:2085, b is an integer of	AL110280,	AC002472, AC004594,	Z98949, AC004686,
			15 to 788, where both a and b	AL022723,	AC006115, AC005488,	AC007298,
			correspond to the positions of	AL021368,	AL080124, AC004690,	AL049759,
			nucleotide residues shown in SEQ ID	AC004808,	AL096776, AL021154,	AL137705,
			NO:2085, and where b is greater	AL021453,	AC004213, AC004159,	AC006112,
			than or equal to a + 14.	AC006039,	AL022336, AL022147	
2086	Н МНН О57	899885	Preferably excluded from the	AI798964,	AA886924, AW082915,	AI015790,
			present invention are one or more	AI888102,	AW305088, AW249524,	AI677907,
			polynucleotides comprising a	AW249655,	AI685359, AI420026,	AW250288,

	nucleotide sequence described by	AW008642,	AI568918,	AW245195.	AI095605.
	the general formula of a-b, where a	AA307509,	AA425494,	AA146920,	AI079724,
	is any integer between 1 to 1336 of	AA742403,	AA628536,	AA425289,	AA393886,
	SEQ ID NO:2086, b is an integer of	AI075449,	AI301574,	AW020330,	AA148122,
	15 to 1350, where both a and b	AA738372,	AA633222,	AI908262,	AA465300,
	correspond to the positions of	AA463585,	AA393791,	R15429, AI55454	IS54546, R16169,
	~	AA629523,	AI193861,	N50479, A	AA234353, AI86383
	, and where b	AA770378,	AI927526,	AA463677,	R24974, AA384622
	than or equal to a + 14.	AI289080,	AA143495,	AA516015,	AI039133,
		AA305089,	AI094204,	AA234408,	AA653256,
		AW026433,	Z45471, Z4	Z41168, AA13	35180, AI541233,
		AA135354,	AI654673,	AA746823,	
		AW337352,	AI907894,	AA152118,	N93532, AI363444
		AA865095,	T24569, Z2	10397, AAO	
		AW189792,	AW170538,	AA906520, AA143494	AA143494,
		AA886922,	AI382046,	D50645, AC	D50645, AC005726, AC004807
		D50646, A7	A74812		
899913	겁 .	Z99396, AW	70,	AL038837, AI	AL037051, AL036725
	present invention are one or more	AL036418,	AA631969,	4	
	ö	AL039564,	AL036858,	AL039156,	AL039108,
	nucleotide sequence described by	AL038509,	AL039109,	AL039128,	AL036924,
	al formula of a-b, wher	AW384394,	AL119484,	AW363220,	AL037094,
	teger between 1 to 702	AL039659,	AL038531,	AL036196,	AL039625,
	:2087, b is an	AL039648,	AL045337,	AW372827,	AL036767,
	15 to 716, where both a and b	AL119457,	AL037082,	AL043003,	AL037526,
	d to the positions of	AL036190,	AL119497,	AL037639,	AL119319,
	de residues s	AL039678,	AL039629,	AL119324,	AL039423,
	, and w	AL036238,	AL038447,	AL039150,	AL119439,
	than or equal to a + 14.		AL119443,	U46350, AL	U46350, AL040992, AL042909
			U46351, AL119483,	119483, AL	AL119363, AL119355
		_	U46341, U4	6349, AL11	.038520,
			AL037726,	AL119335,	AL119418,
		AL039410,	AL038851,	AL039386,	AL119496,
				AL119444,	AL037205,
		AL134530,	AL036998,	AL036733,	AL037615,
		AL134519,	AL134531,	AL119401,	AL134132,

				AT.134527 AT.134528 AT.	AL.043147 1146346 AI.037178
				AL042614,	AL119464
				AL042544,	AL119399, AL042984,
				AL042965, AL042975, AL	AL042542, AL134538,
				AL036765, U46345, AL03	AL036191, AL042989, AL036719,
				AL042551,	
		_		AI142134, AL037021, AL	AL037054, AL036774,
				_	AL036158, AR066494, AR060234,
					A81671, AR064707, AR069079
2088	HTOHV42	510006	Preferably excluded from the	AI014506	
			present invention are one or more		
			polynucleotides comprising a		
			nucleotide sequence described by		
			the general formula of a-b, where a		
			is any integer between 1 to 1410 of		
			SEQ ID NO:2088, b is an integer of		
			correspond to the positions of		
			nucleotide residues shown in SEQ ID		
			NO:2088, and where b is greater		
			than or equal to a + 14.		
5089	HWLX002	900162	Preferably excluded from the	AW373239, AW372628, N2	N27996, AA377857, AA422157,
			present invention are one or more), AW393029,	AA326416,
			polynucleotides comprising a	R54681, AI827898, AI82	
			nucleotide sequence described by	R50597,	AI934499, AW006103, AI422225,
			the general formula of a-b, where a	AI088893,	AI217369,
				AI380811, AI469281, AA	AA494534, AA975272, N21338
			SEQ ID NO:2089, b is an integer of		
			15 to 1226, where both a and b		
			correspond to the positions of		
			nucleotide residues shown in SEQ ID		
			NO:2089, and where b is greater		
			than or equal to a + 14.		
2090	HWLKM7	900249	Preferably excluded from the	AW409927,	
	7		present invention are one or more	, AI979175,	
			polynucleotides comprising a	AA593923, AA573915, AI	AI652793, AI675562,

		nucleotide sequence described by	AT683795	ATGCCGTA	ATOBICAN	3	
		ral formula of a.b. where	, היים היים היים היים היים היים היים היי	ATOTO 12	A1203012,	A1964843,	
		is any integer between 1 to 1618 of	AI380162.	A1656045, A1361395	A1983786,	AI984139,	
		SEQ ID NO:2090, b is an integer of	AA588051.	AT590585	AT673630	A14/2050,	
		15 to 1632, where both a and b	AW206967,	AW137010,	AI288836,	AW170399,	
		correspond to the positions of	AI287323,	AW271527,	AW197398,	AW193824,	
		de residues s	AI380626,	AI869939,	AI371858,	AI650707,	
		, and where b	AI861931,	AI201641,	AW050592,	R00081, TS3	T53389,
		than or equal to a + 14.	AA937517,	AA552662,	AW304869,	AI015077,	
			AI309572,	AI262657,	AI460271,	AI932957,	
			AI950720,	AI652807,	AA327548,	R72802, R50	R50426,
1000			AI634175,	AI089131,	AI986002,		AI659375,
			AI986009,	AI880486,	AI418738,		126655,
			AI719489,	R52030, AA327517, AW272341,	327517, AW	272341, AA5	AA523545.
			AW241543,	AA936966, AI918271, AI652616	AI918271,	AI652616,	•
-			AW197366,	H26610, AI968929,	968929, D2	D25775, AW087283	,283,
			AA100205,	AI880487, D84239,		AC006950, 19574;	742.
			AI479949				
HWMCJ06	900555	$^{\circ}$	N52439, N7	N77401, AA58	AA585439, AI52	AI525556, AI535	5639,
		present invention are one or more	AA585434,	AA585440, AA585453,		316,	,28355,
		g	AI541510,	AI546855, AI525328,			
		nucleotide sequence described by	AI541514,	C15189, AI541523,		AI556967, Z30	Z30131,
		al formula of a-b, where	AI526180,	AI546999,	-	10	
		teger between 1 to 2415	AI541534,		AL045991,	AIS57807,	
-		SEQ ID NO:2091, b is an integer of	AI526140,			AI546828,	
		15 to 2429, where both a and b	AIS41017,	_	_	AI526194, C	C16300,
		d to the positions of	AI546899,		AI541535,	AIS47039,	
		de residues s	AI526196,		AL044029,	AL036500,	
		, and where b	AL134123,	AL043950, 1	AL040252,	AI540967,	
		than or equal to a + 14.	AI535660,		AI541508,	AI541307,	
			AI557262,	AI535813, 1	AI525653,		T11028,
			AL044771,	AL049007, ;	AL043468,		
			AL046147,		AL040768,	AL044377,	
			AI536138,			AL042712,	
			AL043201,	AL040414, 1	AL040571, /	AL046097, D	D61254,
			AI557082,	AL037341, I	R29445, AL	AL079876, AIS	AIS57787

		AL043604, AL044583, R28735, AL048647, AL040510,	AL040625, AL045817, AL041142, AL041238,	AL041133, AL047183, AL040322, AL041131,	AL046330, AL041051, AL041292, AL040119,	AL047170, AL047057, AL04721	AL040463, AL039915, AL04361	AL041197, AL040155, AL041346, AL040529,	AL041096, AL047012, AL041358, AL041277,	AL041163, AL041098, AL040621, AL043538,	AL041324, AI526144, AL040464, AL044162,	AL041086, AL043496, AL041296, AL041233,	AL047593, AL043467, AL041159, AL045725,	AL044186, AL041140, AL040193, AL044037,	AL040091, AL040128, AL040168, AL040255,	AL040285, AL040342, AL040332, AL040617,	AL040553, AL045684, AL040745, AL040370,	2, AL040839	AL040149, D57491, AL043775, AL04165, AL043492,	AL041602, AL045920, AL041278, AL038838,	, AL041635,		, AL040263,	2, AI525320,	, AL041730, AL041523,	, AL046392, AL041374,	, AL043537, AL039338,	, AL039316, AL043923,	, AL043848, AL041459,	, AL044258, AL044201	AL038532, AL037727, T23985, AL040576, AL046914,	AI142134, AI546891, AL045753, AL044274,	AL079878, AL049018, AI557796, AL04044,	AL039744, AL045857, AIS46875, AL038822,	AI525321, AL046327, AI541013, AL041168,	AA585476 AL049069 AL043444 AL041246
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, AL041347,
 , AA585438, T41289, T23957, AISS
, AL080031, AI541345, AL045989, R291
6073, AI557155, AI525203, AI541048,
 AL042096, AI557279,
 6, AL133620, AB03
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 I66486, I66481, A83642,
 I66485, I66490, I66491,
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 5, AR062871, A917
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A43189, A43188, A85395, A85476, A68112, A68104,
A06419, A21892, A23997, A68114, A89633, A89634,
AR057731, AR037157, AR054109, A21895, AR067732,
 AR028564, A05160, A08030, A20502, AR027319,
 A58522, A91751, AR027318
A84772, I19516, A58523, I1
, A22413, A84773, A84775,
, A32111, I63560, AR009152
 , I63563, A60985,
, I08776, I15353, A81878,
, I26929, I44515,
 , I26927, I44516, I18895, E16678,
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, A63954, I91969,
5, A98420
A98427, I6
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8, I08051,
AR031374, AR031375, AR0209
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	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 888 of SEQ ID NO:2092, b is an integer of 15 to 902, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:2092, and where b is greater than or equal to a + 14.
•	969006
	HCRPZ48
	2092

			AL037178, AL042965, AL042975, AL042542,
			AL042989
			9, AL043019, AL042551, AL036191
			AL042450, AI142134,
			AL119464, AL036774,
			AL036836, AL036999, AL036886, AL036158,
			AR066494, AR060234, AR023813, A81671, AR064707,
+	1		
2093 HCKMU04	22006	bly excluded from t	AA258479,
		present invention are one or more	AI337232, AW372227, AI739102, AA505288,
			AI418892, AA551238, AA853934, AI936957, R52096,
		nucleotide sequence described by	AA770298, AW
		the general formula of a-b, where a	AI086723,
		is any integer between 1 to 1801 of	AI094613, AI096869, AI922132
		SEQ ID NO:2093, b is an integer of	
		15 to 1815, where both a and b	AI356823, AA287330, N94480, AA524286, AW005778,
		correspond to the positions of	AW191028,
		רס	AI421557, AI361016, AI359797, AI362874,
		, and w	AI880712,
		than or equal to a + 14.	AA481480, AA291405, N20109, AI263664, AA570059,
			., W94068, AI381877,
			I539565, AA789159,
			Z40719, AA400811,
			5, T58139, AI034063,
			A953460, AW131152, AI146352, AW0549
			AA921717,
			AI083784
			7, R22588, Al
			, AW05783
			φ,
			5783,
+			6, D87444, AL049539
2094 HHBEA82	900784	걶.	\vdash
		present invention are one or more	AI635347, AA195244, AA411217
		polynucleotides comprising a	AI640606,
		nucleotide sequence described by	

		the general formula of a-b, where a	AW081124, AI373594, AW117198, AI424073,
		ny integer between 1 to 5445 of	AA236948,
		:2094, b is an inte	AI041076, AA742216, AA977785, AI979247,
		15 to 5459, where both a and b	AW073726, AA436906, AI129863, AI359758, N24934
		correspond to the positions of	AA491080, AA971157, AI081860, AA490894,
		nucleotide residues shown in SEQ ID	AL135446, AI077569, N32934, AI167862, AI623813
		NO:2094, and where b is greater	
		than or equal to a + 14.	AA906102, AA293745, T27536, N29816, AA640194
			H97513, W73436, AI359073, L44338, AI040170,
			AA931607, AW079283, AI018416, AA235854,
			AA386013, AA307874, H94085, AA782504, AA742947,
			W37849, W69386, AA604174, AI540240, AA805133,
			AI695574, AI537063, AI337935, AA411218,
			AI371459, W73359, AI422480, W74279, R50230,
			R07065, R31685, H94073, AA731784, AA434174,
			AI357532, AI687230, T27535, AA579916, AA588389
			AW103819, W69387, AA101857, AI873792, AI9512
			C00310, R50175, Z24849, AA152394, AI244588,
			AA904357, R67423, AA761110, AA860891, AA935867,
			AI126673, N30780, F00170, D29461, AA377229,
			AA399529,
			AA374839,
			AI913234, AI741350, AI920850, AI018184,
			AA702114, R81654, D29114, AA152500, AA148355
			H94072, N41550, W37848, AF106037, AF222340,
			AF183569, AB011097
2095 HWHGX93	900838	Preferably excluded from the	
		present invention are one or more	, AW385445,
		polynucleotides comprising a	AI337868, AI983250, AW262130, AW337212,
		nucleotide sequence described by	AW305087, AI587497, AI826854, AI640371,
		al formula of a-b, where	
		teger between 1 to 207	AA927991, AA071469, AW373440, AA513750,
		SEQ ID NO:2095, b is an integer of	, AI696797, AA922948,
		where both a and	, AI920995, AI624419, W92531, AI
		correspond to the positions of	AI828286, AI379231, AI091871, AI584063, W7222

0, AI818524, AI378538,
AI674870, AA449300, AI925019, AA431859,
AI608680, AI435229, AI627567, AI587133,
8, AI354309,
4, AI084022, AA44974
AA431858, AI366084, AA505877, W77968, AI911667,
6, N64004, AA976403, AW337258, N32415,
2, AW136886, AI12403(
4, AW193263, AI431982
AI952361, AI223458, F37472, AI401365, AI290429,
AW058441,
2, AI950830, AW276587, AI35732
, AI220027,
I453327, AW
, AI950575, AW316754, AW304759, H1612
AA024771, AA335712,
N32424, A
N56835, AA602994, D79675, AI537354,
AA347786, D62623, AW263293, D62595
D62097,
8, D62131, D79867, AA371169
, AW263466, D62031,
, AI802265, D61938, AI28000
, D63012, De
i, AA082155, AA297695, C16543
37, AI686490, D62783, H877
AA297550, C02046, AA095691, H2
1, D79835, AA329099
3, AW021588,
, AI637584, AI498067
AL039086, AL121496, AI281772, AW169671,
344,
AI362637, AL045266, AI476046, AW088134,
AI933589, AW190042, AI922676, AW088903,

AI92	1248, AI289629,
AIS	
AW3	AW302988, AW103371, AW073994, AI537677,
AI9	AI909697, AI269862, AI868831, AI922901,
AIS	, AI524671, AI88446
AI8	6, AI670009, AI802542,
AW0	AW081298, AI783504, AW268122, AI625701, U8856 [.]
AF017	D50462, AF017986,
148	89947, AF113013,
AFC	A08913,
AL1	AL133093, A08910, AL133080, A77033, A77035,
AFI	i, AL122050, AL133075, A08909
145 J	82022, AL117435, AF113019, AL
ALC	AJ238278, AF104032, Y11587,
ALC	\sim
Y16	Y16645, AL137557, AF017152, AF177401, AL050393
ALC	AL049382,
ALI	AL110225, A65341, AL137271, AL117460, AF090903
ALC	AL080124, AF078844, AF118094, AL133640,
ALI	AL110221, A12297, X65873, AL122121, AL137550,
878	, AL117583, AL133557
ALI	, X98834,
Y11	
AF1	AF113694, AF113691, AF153205, AL049466,
AJC	AJ000937, AF113699, X84990, AF090900, AF146568,
ALC	0, I03321, AF091084, X8
ALC	4, AF118070,
ALI	AL122098, AL117457, AL050116, AL133016,
AF1	AL050108
ALC	AL080127, X72889, U35846, A58524, A58523,
E03	158248, AL
AF	AF113690, AL049464, U80742, A08912, AF113689,
AFI	AF118064, AR059958, AF113676, AL133565, X6357
AL:	4
ALI	96, AL096744, AL050146,
AL	AL133606, L31397, X96540, AL122110, E07361,

9, R00450, AI	T04952, AI654393,
D30892, AA	Ā
	AW439050, AW374583,
AA927954, AW374501, AW374	1602,
 AW051233, AW152182, AI34	5347,
AW169624, AW050578, AW020	419,
AI612750, AI915291, AI872	72555, AI678395,
5,	9828,
AI673785, AW059828, AIS:	3885
AI890907, AI689388, AWI	89415, ALO37454,
8, AA9	2853
AW088697, AA502794, AI3	AI345608, AW117919,
AI538885, AW088628, AI5	AI521799, AW020693,
AI340603,	AI540759, AI345396,
AI345471, AI538055, AI6	AI628833, AI251221,
AL039086, AI249257, AA4	AA420722, AI345745,
AI287477, AI540674, AI4	AI446373, AL038605,
AI570912, AI284131, AI824	24576, AI690748,
, AA176980,	AI784219, AI242248,
AI340519, AI355779, AI3	34714, AI434453,
669, AI922215,	3559, AI783
6, AW161579,	35363,
AI281888, AI284517, AI9	
	AI860027, AI553645,
7, AI889372,	0013,
88, AI623941,	4201,
4, AI473536,	4153,
AI799195, AW151766, AI69	8391,
 AW022699, AI554343, AI3	AI311892, AI475139,
AI434464, AW162194, AI28	80655, AI345415,
 AW020095, AW074869, AI88	9189,
AA641818, AI868204, AA8	AA848053, AI633125,
	AW080079, AI538564,
45, AI579991,	5200,
63834, AI859464, AI6	3859, AF02079
AF067146, Z83841, I4897	8, A93350, I89947,

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				AF114170, AL122049, I09360, A65341, AL050024,
				, AL117460, AL117585, AL080158
				AL117578, AL137712, AF026008, AL133077,
		_		AF106657,
				92, AJ00
				D83032, S78214, E07361, S61953, Y10080, S83456,
				1, AF097996, Y115
	-			
				7,
		_		~
				U67958, X06146
2097	HCRP045	996006	Preferably excluded from the	, AI207822,
			present invention are one or more	1, AI951947,
			polynucleotides comprising a	AI494358, AI262716, AW245429, AW341614,
			nucleotide sequence described by	AW378336, AA894688, AI801517, AI076235,
			the general formula of a-b, where a	AI828126, AA904279, AI675937, AI460342,
			is any integer between 1 to 3081 of	AI128285, AW294715, AI745086, AI077325,
			SEQ ID NO:2097, b is an integer of	AL042068, AA115771, AI624138, AI082386,
			15 to 3095, where both a and b	AA459947, AW167502, AIS40099, AI635602,
			correspond to the positions of	AI364603, W67744, AW341954, AW078482, AI434372,
			e residue	AI953308, AW261951, AW005837, AI633304,
			NO:2097, and where b is greater	AI368673,
			equal to a + 14.	
				N23827, AI
		_		
				7, AI334786, AA641824, H14234,
				\vdash
				7, AW082444, AI363715,
				AW190481, AI439839, AI032415, AA552221,
				AI825222, AI473352, AW378284, AI198651,
				AA729339, AI784394
				F28183, AW370330, AI870988, H69297, H28474,
				AI356300, H58452, AA113878,

			AA470038, AA115770, D54636, AW263948, W92582.
			H69392, C17841, C17347, AA934363,
-			8, R54975, AI669735, AI6903
-			, W32110,
			A968922, AI351193,
			H58453, AA421734,
			AA296340
			AI669906, T29492,
	_		
			AW378303, AI279499, AW372174, AI560558, C17284,
			1587988, AI684276, H65933,
			AA581394, R62494, AI696415, AI299631, AW378274,
	-		H39968, AA460036, AA641823, AI964054, D20567,
			8, AI479661, AW237023, AL117507
			X04654, X07401, AL117399, X84841, M57939,
			X06814,
			M57936, X07402, X07403,
+			
2030 HWLWF0U	166006	乙 :	AW368386, AW238539, AA029705, AI831658.
		present invention are one or more	AI880448, AI826080, AI80944
		ides comp	, AA21659
		deg	1, AI753535, AA425993,
			0, AA405525, AA293346, AL03803
		teger between 1 to 1400	5, AI831455,
		SEC ID NO:2098, D is an integer of	5, AW102926, AI609085,
		13 to 1414, where both a and b	AA449167,
		d to the positions of	AW168852,
		de residues show	
	_	, and where b	AI148267, AW118509,
		than or equal to a + 14.	AA826234, AI144475, AW16985
			AA425864, AI075654, AI01848
			AI023124, AA457092, AA57750
			AA229090, W00840, AI199803, AA405536, W45253,

·	AR85101, W89773, W80519, AA926675, AA643707, AA85101, W89773, W80519, AA926675, AA643707, AA85101, W89773, W80519, AA305369, AA76469, AA130762, AA9306169, AW39252, AI744974, AA130762, AA930180, AW39252, AA06545, AI148913, A166661, A1126691, AA705545, AA140947, AA877983, W30924, W62547, AA570576, AA06545, AI148913, W30924, W62547, AA570576, AA06546, T2973, AA399426, AR33884, W42994, AA05466, T2973, AA599426, AR38849, W32994, AA0512, W30287, AI132789, AI182584, AA253113, W69812, W30287, AI132789, AI182584, AA868114, AA00747, AI132369, AI122584, AA868114, AA007917, AA10747, AI132789, AI831831, AA902522, T09075, AA675190, AL038014, AI831831, AA902522, T09075, AA577290, AL038014, AI83184, AA19886, AI470710, AA502591, R391371, AA19886, AI470710, AA502591, R391371, AA198876, AI4407, AI4407, AA85386, R49107, AA300055, AA4381377, AA464289, H44177, AI085238, C03394, AA454676, T35412, AA459695, AA533748, H94767, AI18861, AA706052, H04146, AA333748, H94767, AI18861, AA706052, H04146, AA333748, H94767, AA16813, AA706052, H04146, AA333748, AA706615, AA706052, H04146, AA365396, RW381377, H42139, AA403384, AA4035896, RW38137, AA706052, AA403384, AA41657, AA866125, AA41335, AA366180, AA4165, AA195897, AA866125, AW367180, AA20986, H04145, AW195897, AA866125, AW367180, AA20986, H04145, AW195897, AA866125, AW367180, AA20986, AA1187, AN364896, AA866125, AW467866, AA409386, AA1187, AA195897, AA866125, AW367180, AA209868, AA1187, AA195897, AA866125, AW367180, AA209862, AA209868, W31187, AA209818, AA1103, AW364866, AU44185, AW195897, AA866125, AW367180, AA209862, AA209862, AA11872, AA1187219, AA11874, AA709865, AA4099866, AA11872, AA1989897, AA866125, AW668999, AW444466, AA2098868, W31187, AA866125, AW469866, AW469866, AA1187, AW195897, AA866125, AW4698969, AA4099868, W31187, AA7098699, AA809886, AA1187, AW3698997, AA866125, AW4698969, AA4099868, W31187, AW369899, AW4998868, W31187, AW389899, AW4998869, AW389889, AW389899, AW389899, AW499889888, W31187, AW389899, AW4998898, W31187, AW389899, AW49988989, AW49988888, W31187, AW389899, A
	AA977681, AA303937, AA357438, 79215, AA654730, AA654815, AA42 A164587, AA658950, AA907646, AA
	4, N90989, AI952123, AA523768, AA428055, AI768287, H24905, AW

AA844041, AA527916, AI783869, Z46157, AA1	AA126575,
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AF151812, AF091085, M30773, AL137521, X63	410,
I48978, X66871, AL117460, U75932, AR068753	3,
 AL049382, AL137548, AL137476, A15345, U58996	966,
6, A08910,	226,
 AF100931,	137,
	947,
 A08907, M30514, A08913, AR038854, AF113677	7,
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 U49908,	
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 AL133016,	
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	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 2157 of SEQ ID NO:2099, b is an integer of 15 to 2171, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:2099, and where b is greater than or equal to a + 14.
	900993
	HCNCY 58
	2099

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AL039625, AL039648, AI557807, AL045337,
 AL036924, AI526140, D29033, AL037094, T23888,
 AL039629, AL039150, AL039423
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AJ238010
Z32836, I84553, A92133,
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AR038762, A11245, A02135, A04663
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 A77094, A
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D17247, A93923, Y11449, I01995,
AR008429, A93916, Y11447, I25027, I26929,

2100 F	HCNDA61	901111	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 1172 of SEQ ID NO:2100, b is an integer of 15 to 1186, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:2100, and where b is greater than or equal to a + 14. Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 3095 of SEQ ID NO:2101, b is an integer of 15 to 3109, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:2101, and where b is greater than or equal to a + 14.	126928, A98420, A98423, I26930, I2693, A98436, A98417, A98427, AJ231028, A98436, A98417, A98427, AJ231028, A98436, A98417, AR069079, AR051957, D50010, A11458, A22734, E17098, AB026436, AL133053, AL122101, I19525, AL133074, AL133053, AL122101, I19525, AL133074, AL133053, AL1320845 10, A06631, AJ230845 11, A184054, AA634246, AA630382, AR178852, AI825946, AA630382, AI6232766, AA931283, T24595, AI6232766, AN369427, AN176607, AN176607, AN1888177, AA992910, AF061022, AF06133, AN361521, AN361521, AN361521, AN361521, AN361522, AN009764, AI687981, B, AA305409, AA514186, D80166, D58246, AI53566, D80439, D81026, D51221, H67854, D80022, D81030, D81111, D8013 D80212, D59619, D80251, D59783, H67866, D59859, D59551, D80165, D80268, D80366, D80251, D80251, D80268, D80366, D59889, D80168, D80268, D80366, D59889, D80168, D80268, D80366, D59889, D80168, D80268, D80366, D59889, D80166, D80251, D60251, D60251, D60251, D60251, D60251, D60251, D60268, D80366, D59889, D80366, D80268, D80366, D59889, D80366, D80268, D80366, D59889, D80366, D80268, D80366, D59889, D80366, D50268, D80366, D59889, D80268, D80366, D50268, D80366, D59889, D80366, D50268, D80366, D59889, D80268, D80366, D50268, D60268, D6026
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	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 1424 of SEQ ID NO:2102, b is an integer of 15 to 1438, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:2102, and where b is greater than or equal to a + 14.
	901128
	HWLRB02
·	2102

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AL12212
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AC00425
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AC005548
AC005821,
AF042484,
AP000500,
AC002525,
AL117667,
AC008170,
AC005632,
AP000475,
AL022400,
AL034371,
AC00011
AL035467,
AL079352,
AC007156,
R40742,
H86543,
W96135,
AA021123
AA035574
AA102640
AA156568
AA236317,
AA25880
AA43050
AA513740
AA568471
AA568809
AA57699

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AW132045,	AI688692	
AI289782, AW132045.	AI687053	
AA233673,	AA723802	
AI860695,	AW167654	
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14782, T09341,	13, T74334, AI74	
AA78889	51, H29344, AA77	
AW243083,		
ν ω α	A1610646 A1635849	
		1,
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AI582535,		
AI559582,	6	
AI417451,		
AI380398,		
AI301333,		
AI280684,	AI275808	_
114, AI261269,	3, Z41712, F073	STOTE PECEFCIA
AA909244		
AA670172,	AA669335	
AA411709,	δ,	
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AA977546,		
AA923484,	3,	
AA030222,	ສັຕ	

			15 to 2519, where both a and b	AT337294	ATREACTE	ATREJETE	10000000	
			correspond to the positions of	AI652837,	AC005837,		X11274 A59344 AL10209	50000
			nucleotide residues shown in SEQ ID				11100	7
			NO:2104, and where b is greater	-				_
			than or equal to a + 14.					
2105	HPBEM10	901276	Д	AA287703,	AA287702,	AA365652,	AA282618,	
			present invention are one or more	AA927786,	AW364617,	AA027167,	F24601, AJ	AI968421.
			polynucleotides comprising a	AI913352,	AI302397,	AI040349,		AA355129
			nucleotide sequence described by	AI984941,	AI184494,	AA480189.		, ,
			the general formula of a-b, where a	AA027168,	AA382209,	AI935351,	AB023172	•
			is any integer between 1 to 1298 of				1	
			SEQ ID NO:2105, b is an integer of					
			15 to 1312, where both a and b					
			correspond to the positions of					
			nucleotide residues shown in SEQ ID					
			NO:2105, and where b is greater					
			equal to a + 14.					
2106	HWBDL33	901333	Preferably excluded from the	AI263085,	AI671224,	AI741604.	AW055187	H93009
			present invention are one or more	AW057512,	AA058688,	AI800594	AW195361	
			polynucleotides comprising a	AI740946.	AW271301.	AW292805	AA160279	
			nucleotide sequence described by	AI302809.	AA160278	AT769897	AT200257	
			the general formula of a-h where a	AT628787) T T 2 E 2 7 2	71,000,1,	A120021,	
				, 10201A	A1/332/3,	A1458862,	A1091306,	
			ceder Derween I to 185	AWZ/2744,	AI128201,	AA716336,	AI707638,	
			15 to 100:2106, D IS an integer of	AA031623,		N59386, AA421911,		AW052091,
			is to 18/1, where both a and b	AA088175,	AI824017,		AA461046,	
				AI635515,	_	AI699923,	AI880867,	
	-	-	de residues show	AI597746,	AA460478,	W03796, AI239461,	239461, AI	AI863568,
				AA448335,	AA582895,	AA449267, AI278475	AI278475,	•
			than or equal to a + 14.	AI691016,	AI758904,	H64963, AI	H64963, AI278932, AA709030,	709030,
				AI418284,	AI361585,	AA045175,	AA150151,	
-				AI634797,	AA035209,			H59637,
		_		AA035208,		AA917066,		
				AI300367,		T97469, AA	T97469, AA502528, AI199994	199994,
				AA974453,	AA810540,	AA411404,	AA411404, AA576365, F20467,	F20467,
				AA040431,	H	373386, AI	684553, AI	AI962642,
				AI474422,	AW072561,	AI824266,		N73170,

				AA731356, A	AI806247, T97468, AA502505, H13072,
					H64964, T96890, T96889, R58859,
				AI161128, A	, H95741, AA3802
				T70436, H94	T70436, H94235, AI305839, AA366448, AI743473,
				AI668883, A	AA366209, R97096, AA502417, T81549,
				AA361023, A	AA045294, AA976534, AA974771,
					AI922795, AA441989, AW148422,
				AW182457, F	AW182457, H13276, AA344621, N77074, AA713812,
				W01926, AA031704,	131704, AI733416, AA736644, AA040430,
				AA101990, N	AA781193,
				AW452710, A	AA152220
2107	H2LBA47	901375	Preferably excluded from the	AI346914, A	AW361114, AA573910, AA573949,
			present invention are one or more	AA314779, A	AA573904, AA573811, AA573823,
			polynucleotides comprising a	AI791286, A	AI791498, AA573762, AA308533,
			nucleotide sequence described by	AI732541, A	AA314573, AA315990, AA307789,
			al	AA308019, A	AW362522, AA315862, AI925615,
			is any integer between 1 to 1295 of	AI802703, A	AA315993, AA313200, AA316848,
			SEQ ID NO:2107, b is an integer of	AA316249, A	AA552253, AA316525, AA552098,
			15 to 1309, where both a and b	AI393251, A	AI926615, AA313549, AA508861,
			correspond to the positions of	AA316634, A	AA552332, AA552296, AA314847,
			nucleotide residues shown in SEQ ID	AA573769, A	, AA315069,
			NO:2107, and where b is greater	AW130226, A	AA552106, AW363214, AA552304,
			equal to a + 14.	AA551912, A	AA316658, AA552492, AA574080,
				AA314181, A	AA552602, AA307590, AA315842,
				AA552328, A	AI888532, AA588112, AI318255,
		_		AI318551, A	AW362532, AI307602, AI452604,
				AA551820, A	AA315757, AA313418, AW351498,
				AW361505, A	AA584947, U54601, AW130541, AW182560,
				AA612996, A	AI691058, AI933755, AA527185,
				AA588123, A	AA316515, AI537454, AA581266,
			-	AI282560, A	AA583270, U54606, AA582738, AA535703,
				AW351551, A	AI732344, AA837983, AW361468,
				AI470732, P	AW044042, AI444965, AI652625,
					AI919553, AW008048, AL036638, N71180,
				AW020397, N	N75771, AW020710, AW409775, AI557238,
				AI932458, P	AI698391, AW029401, AI818358,

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	AF078844, AL1	137478, E06743, AL137574, AF061795,
	AF151685, S76	S76508, U88966, AF120268, AF113676,
	AJ012755, I89	AL023657, AL110218, E1274
	Æ	I49625, A08907, AF
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	A58523, A21103,	Y10655,
	I33392, X80340,	10, E02349, AJ238278, AF094480,

E08631, AL080140, AL137521, AF026816, U75932, S61953, S75997, AR020905, X82434, E03348, AL050024, AL122050, AL080159, AL133640, AF183393, X52128, U00686, AL096744, I66342, AF040751, AL137533, AF061981, U80742, I32738, U72621, AL080148, AL080126, AB008792, AL137292, AF104032, X56039, AB008791, I41145, X66862, AL049339, AF113699, AR029490, Z82022, AF162270, AL133557, AL080127, AL110221, AF090900, Y08769, AL1325093, A23630, AL110222, AL133112, X96540, A08911, A18788, AL110159, AL049300, AR038969, AL137560, AL117583, A21101, AL133665, AF090896, AL133560, AL117648, AF067790, AL110280,	AI634717, AW369331, AW166169, AW152548, AI080640, AI678847, AI921153, AI828325, AI828325, AI559391, AI025266, AI800451, AI720013, AA316874, AA316874, AA316233, AA316233, AA316233, AA516596, AI610106,
E08631, AL080140, AL137521, AF026816, U S61953, S75997, AR020905, X82434, E0334 AL050024, AL122050, AL080159, AL133640, AF183393, X52128, U00686, AL096744, I66 AF040751, AL137533, AF061981, U80742, I GVZ621, AL080148, AL080126, AB008792, A AF104032, X56039, AB008791, I41145, X66 AL049339, AF113699, AR029490, Z82022, A AL133557, AL080127, AL110221, AF090900, AL122093, A23630, AL110222, AL133112, X A08911, A18788, AL110159, AL049300, AR0 AL137560, AL117583, A21101, AL133665, A AL133560, AL117648, AF067790, AL110280,	AI815198, AI817063, AI817063, AA909945, AA316115, AW364225, AW3164225, AW315049, AA315049, AA315049, AA315629, AA31572, AA314372, AA314372, AA316967, AA316967, AA316967, AA316967, AA316967, AA316967, AA316967, AA316967,
ALO80140, AL S75997, AR02 4, AL122050, 3, X52128, UC 1, AL137533, AL080148, AL 2, X56039, AE 9, AF113699, 7, AL080127, 3, A23630, AL A18788, ALL11 A18788, ALL11 A18788, ALL11 A18788, ALL11 A18788, ALL11	
E08631, Al S61953, S' AL050024, AF183393, AF040751, U72621, Al AF104032, AL049339, AL133557, AL133557, AL133560, AF137367, AL133560,	AI675865, AI888294, AW194118, AA573742, AW190856, AA582017, AA314225, AW272720, AW272720, AW314225, AW314225, AW314225, AW314225, AM582851, AA582851, AA582851, AA582670, AA552670, AA552670, AA552670, AA552670, AA552670, AA552670, AA552670, AA5626, AA314146, AA314146, AA314146, AA314146, AA314146, AA314146, AA314146, AA314146, AA314146, AA314146,
	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 929 of SEQ ID NO:2108, b is an integer of 15 to 943, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:2108, and where b is greater than or equal to a + 14.
	901415
	HCQAJ72
	2108

				AI146786, AA315166, AI582452, AI916480,
				AA593818, AA313235, AA421562, AI285429,
				AI924498, AA583091, AI358508, AA425142,
	-			AI445130, AI888732, AA315613, AA244356,
				AA632103, AW190915, AA581848, AW152169,
	-			AW191880, AI678427, AA565444, AW192785,
				AI891014, AW370274, AA314206, AA476675,
				A1473553, AA625485, AA687567, AI675714,
	•			AA316508, AI685830, AA314052, AI434099,
				AA298537, AI469613, AI972701, AI972499,
				AA526975, AI933636, T86663, AI623264, AA513297,
	-			AIS81525, AW080588, AA501945, AI400863,
	•			AA315408, AA298527, AA639696, AA421527,
				AA558986, AA570785, AW303846, AI537212,
				AI926128, AI695291, AI986354, AA055880,
				AI445127, AW196067, AI580982, AI932444,
				AI919084, T24475, AI471336, AI783818, AI924494
				AA306967, AI867585, T24892, AA506763, AA307841
				AF088867, AF038451, AF007791, AF044262, AB0165
2109	HETHC61	901421	Preferably excluded from the	7, AI492171
			present invention are one or more	AA831769, R25716, AA359492, AW238299, R62460,
,			polynucleotides comprising a	AW379689, C02578, AW241754, AW243207, AI034221
			nucleotide sequence described by	
			is any integer between 1 to 1363 of	
			SEQ ID NO:2109, b is an integer of	
			15 to 1377, where both a and b	
			correspond to the positions of	
		_	nucleotide residues shown in SEQ ID	
			NO:2109, and where b is greater	
			than or equal to a + 14.	
2110	HTXLJ25	901472	Preferably excluded from the	AI829099, N25625, AI126506, AI200037, AI128843
			present invention are one or more	N34223, AA743134, AW024969, N36303, AI217597,
			polynucleotides comprising a	AA605122, AA729493, AI160533, AW450603,
			nucleotide sequence described by	AA568193, AA568681, AW020616, AI695490, N26904,
-			the general formula of a-b, where a	N24885, W52651, AA648514, AA806507, N35103,

			is any integer between 1 to 774 of	N72137, A	AI802647, AI3	AI312534, AA729125	A729125 N34254	Г
			SEO ID NO:2110, b is an integer of	219599	Н86994, Н86	H86995, N39)	
			15 to 788, where both a and b		I032141, W00	385, AW2	AI032141, W00385, AW298649, AA296449,	
				N28403, R	~	1, R2630	4, AW452862,	
			de residues show	AW453038,		A988539,	AI299683, AA988539, W52017, AI039557,	
			, and where b	AI141901,		W236299,	AI361669,	
			than or equal to a + 14.	AI674252,		52444, N	T25829, AI452444, N20053, AW074182,	
_				AI984739,		AA543074,	T25828, AA358828,	
				AA653691,	AI362330,	A1906328,		
	20111011			AL050024,	E03671			
1117	HCINAI22	901473	Ω	AW001287,	AW300770, A	AI691072,	AI936111.	Т
			present invention are one or more	AA622758,	AI245950, A	AA563933,	AA622120,	_
				AI801582,	AI348065, A	AA552519,	AW001308,	
			nucleotide sequence described by	AA847242,	AA622570, AA	AA552362,	AI660557,	
				AW050790,	AA582787, AV	AW000826,	AA643708	_
				AA298484,	AI732367,	AA643616.	AA514424	
				AI673534,	AA857546, AA	AA025434,	AA543029,	
_			15 to 1019, where both a and b	AI821215,	AA470683, AI	AI732198,	AA297147.	
			correspond to the positions of	AI582013,	AA297176, AA	AA025433,	AI749731.	
			nucleotide residues shown in SEQ ID	AA594300,				
			NO:2111, and where b is greater					
9::0			qual to a + 14.					
7117	HSIAL77	901494	ヿ	AI685117,	AA583424, AA	AA554005,	AI718759.	
			present invention are one or more	AI721245,	AI732444, AI	AI832388,	AI732445,	
7			polynucleotides comprising a	AI720621,	AI720903, AA	AA130541,	AI460276,	
			nucleotide sequence described by	AI990978,	AI990957, AA	AA574028,	AI879881,	
•			al iormula of a-b, wher	AI733759,	AA115664, AI	AI832502,	AI983398,	_
_			teger between 1 to 961	AI733760,	AA580320, AA	AA130579,	AA134398,	
			SEQ ID NO:2112, b is an integer of	AA126912,	AA132736, AI	AI748949,	AA308497,	
			15 to 975, where both a and b	AA134332,	AA055636, AA	AA133748,	AA134372,	
			sitions of	AA436898,	AI708072, AA	AA130459,	AA603658,	
			ס	AA134397,	AW204007, AA	AA297640,	AA102277,	
			12, and where b	AI302569,	AA316534, AA	AA130403,	AI983618,	
			than or equal to a + 14.	AA296956,		AA506416,	AI445264,	
				AI688106,		AA100297,	AI963380,	
				AI925567,	AW362172, AI	AI672950,	AW362167,	

AA29852
AA297149
AA297152,
AA297184,
AW365047,
AI581967,
AA298926,
AW028870,
AA1328
AA297182,
AI88039
AA054072,
AI459944
AW189415
A1680162,
AW082594,
AI824576,
AI269696,
AI811785
AI345253
AW168373
AI34885
A1680498
AW022682
AI47812
AI80013
AI81857
AI613471,
AW151785
AI400725
AI872423,
AI493576
AI814087,
AI34573
AI34561
AW08793

	0, AI500077, AI345415,
	4, AI334884,
	AI870192,
	0, AI349967, AI539847, AW08027
	5, AI366985, AI345787, AW10545
	3, AI783504, AI610799, AW30299
	809, AI348897, AI352497, AI92290
	3, AL036631, AW198075, AW088
	3, AI587606, AI783861, AI46887
	9, AW104196, AI611810, AI59012
	9, AIS89947, AI349957, AW14922
	, AA848053, AI924686, AF01483
	1, U82953, X79303, AF091738, U67958,
-	AF036941, AL080127, AF061943,
	`
	AF158248, AF113694, AR000496, U3965
	AL117440, AL137527, AF017437,
	F113689, AF
	AL049314, AJ238278, X84990, AL117585, AL117457,
	AL096744, AL117435, A08910, I89931, A08909,
	I49625, AF003737, AF113019, AL1375
	4, E03348, Y11587, AR059958, I03321,
	3, A77033, A77035, X82434, AB
	9, AR038854, L19437
	, AF067728, AF090903
	AL050393,
	AF113690,
	AL080124, AL137550, AF111851, AF153205, S78214,
	A03736, U72620, AL080060, A653
	, AL1336
	AL122098, AF061795, Y14314, AF1516
	7, AL133565, AL137476,
	1, AR011880, AL122110,
	1, AF113699,
	4, AL11743
	AF061573, S61953, AL133067, AL137478, AL049382,

				U49908, AL080086, AF078844, I26207, AF119337,
				I42402, Z37987, AF090901, X65873, AF079765,
				AL122049, AL137526, AF118064, I09360, X87582,
	-			A
				AL133075, AF113676, AL133077, S68736, AL133568,
		_		AL133014, U80742, U78525, AL133113, E02221,
				,049452,
		~		
				AF057300, AF057299, AL110280, X72889, A58524,
				ᄀ
				AL049464, AF008439, AF118094, AF097996, A90832,
				AL050024, Y11254, AJ000937, AL049430, U58996,
		_		AL137459, AF111849, A93350, Y09972, AL050108,
				AF177401, I00734, AF090896, AL137273, AL122093,
				Y07905, AL133072, U42766, AL137521, AB019565,
				AF067790, AL133104, AL137557, AL049283, X70685
	•			8, AF079763, AJ2428
				E08631, AF125949, L31396, E00617, E00717,
				E00778, U68387, AL050146, AL110225, AL117394,
				A12297, AL133606, L31397
2113	HRACJ32	901515	Preferably excluded from the	
			present invention are one or more	AI377209, AA031514, AI565078
			polynucleotides comprising a	
			nucleotide sequence described by	AA987674, AA908398, AI679314, AA908479,
			the general formula of a-b, where a	AA828906, W31903, W60256, AA528246, W39266,
			is any integer between 1 to 1159 of	AA977868, AI570763, AA970839, AI920871,
_			SEQ ID NO:2113, b is an integer of	AW338549, AI696789, AI962006, AA344350,
_			15 to 1173, where both a and b	AA299556, AA910725, AI219260, AA299411,
			correspond to the positions of	AI921665, AA031513, AA887197, AI888609,
			nucleotide residues shown in SEQ ID	AI925329, AI888421,
			NO:2113, and where b is greater	AA034355, AI537808, AW297694, AA029323,
			than or equal to a + 14.	AA173929, T27577, AI869462, AA335005, AI933599

	R36271,	AW162194, A	AW167918, A	AI559752. AL036638
	3	3	υ C	, ,
	38	AW163823	AI537677	
	AW161156	, AW163554,	37187	939
	AI564290	, AI282930,	AI697324,	ເຫ
	AI554821	, AW161579,	AI687295,	966
-	~	, AA641818,	σ	~
	AW079572	, AI114703,	AI699143,	AA420722,
	AI890223	AL043345,	AI570966,	AI469505,
	₩.	AI802542,	AL036901,	AL037454,
	AI468872,	AI277008,	AL119836,	AI919593,
	AI340603,	AI590043,	AI909697,	
	AI539800,	AW022682,	AI538850,	AI568138,
	AI884318,	AI345416,	AI802240,	AI345612,
	AL120700,	AI698391,	AL042191,	AI345415,
	AI491710,	AI588892,	AI690748,	Z99428, AI683395,
-	AA640779,	AI097643,	AW198112,	, 1
	AA572872,	AI868740,	AI798456,	AI564259,
	AI690411,	AI686576,	N29277, AI	538764,
	AI862135,	AI932638,	AI499285,	~
	AL119863,	AI270295,	AW303089,	AI923989,
	AA580663,	AW169604,	AI862144,	AW051088,
	AW083750,	AI538885,	AL134259,	AI633196,
	9	AL039086,	AI623682,	AW023338,
	AL043355,	AW103628,	AW162071,	AI580436,
	AI624963,	AI934011,	AL119748,	AW088899,
		AL119399,	AI434242,	AI251221,
	AI363957,	AI916419,	AA833760,	AW020693,
	91	AI281653,	AI281867,	AI440263,
	AI473536,		AI475371,	AI624943,
	901	AI80046	AI270055,	AL036274,
	AI954080,	AA572758,	AI824746,	AI241923,
	AL036802,	AL046618,	AI312428,	AL121328,
	AW403717,	AI349645,	AW074869,	AI280561,
	567	AA916133,	AI890907,	AI917963,
	AL036631,	AW059713,	AW150308,	AI570807,

	AI567582, AI863382, AI636588, AI648458,	0,302, A1033321, A1330303, A149430	2, AIGIZ/13, A130/203, 7 AIGEN892 AIK20517	1. AI633477. AW265004. AI597	3, AI567866, AI827440, AW08957	9, AI699865, AW024564, F27788,	1, AW028840, AA693347,	AI969655, AI950664, AI340519, AA908294,	7, AI624293,	AI634736, AI638798, AW051059, AI690813,	, AI812015,	191443, AB031324, AB031323, L24374, L22524,	1,22520, X63162, 1,22523, 1,22521, 1,22522, 1,22519,	X07821, X80340, AF039138, AF039137, AL050116,	AL122098, AL133081, I89947, I48978, AF057300,	AL137271,	AL133067,	AL137459, AB007812, I03321, AL049382, AB016226	580, AF1136	AL137527, AL122123,	AF100931, AR038969,	AL137557,	AL13364	AL137558, AL137488, E02221, S76508, I89931,	8912, AL133606, MB	1137283, AL049300,		110, S759	X57961, I68732, S61953, A65341, AL133080,	110197, %84990,	AL117435, I00734, A08910, U49908, AF090934,	15822, AL137550, AF090880			AL137521, E12747, D83032, AL133560, AR038854,
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901	AF008439, A21103, AF078844, AF113677, AF118094, AF159615, Z37987, AL050149, AL133016, AL096744, AF158248, S68736, A15345, AF113019, X82434, AL049430, AF125949, AF177401, AL117432, AF113691, AL137480, AL080163, AF032666, AL137479, AF126247, X79812, AF118070, AL137640, AJ242859, AL122100, AF061795, AF151685, AF106862, I49625, AF017437, AL049452, AF176651, AF090900, X98834, I89934, AL080086, AJ000937, AF113690, A18777, AF097996, AL06452	AW340394, AW245451 AW340394, AW245809 AI832220, AI376745 AI917768, AI536948 AW118765, AI751172 AI635792, AI480259 AI992041, AI2117673 AW002588, AI360270 R56232, AI631567, A884481, AA354601, W276496, C00611, R9 W02478, AA9695594, AA616840, R96719,	Preferably excluded from the present invention are one or more polynucleotides comprising a nucleotide sequence described by the general formula of a-b, where a is any integer between 1 to 1863 of SEQ ID NO:2115, b is an integer of 15 to 1877, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:2115, and where b is greater
900.			

			than or equal to a + 14.	
2116	HSSGC06	901621	Preferably excluded from the	AA612669, AW026486, AA612668, AI458253,
			present invention are one or more	AA311709, AI859961, AA005340, AA005433,
			polynucleotides comprising a	AA397884, AI751088, AA005434, AA932249,
	-		nucleotide sequence described by	AW273329, AA287706, AI016843, N66090, AI205137,
	-		the general formula of a-b, where a	AA488248, W90552, AA699684, AI694508, W90553,
			is any integer between 1 to 814 of	AA130969, AA609505, AA399646, AI693778,
			SEQ ID NO:2116, b is an integer of	AA099841, AI201786, AI452981, AA644003,
			15 to 828, where both a and b	AI085190, AI808813, AI202524, N98636, T60671,
			correspond to the positions of	AW407236, R09367, AA191378, AA827388, AI276380,
			nucleotide residues shown in SEQ ID	AA488193, H23331, AA160239, AA309096, F12355,
			NO:2116, and where b is greater	AI142701, T57771, AA085583, T64868, AA310662,
			than or equal to a + 14.	AA357288, D58848, AA055733, R09250, AI183865,
				AA356179, M78761, AA045074, AA461214, AA190768,
- ,,,-				T80323, AW363425, AI677821, R17951, AL031685,
				AF131742, AA827467
2117	HSICN14	901875	Preferably excluded from the	AL120519, AL120518, AW167654, AI860695,
			present invention are one or more	AW340140, AA878120, AA824284, AI829215,
			polynucleotides comprising a	AI858970, AI983809, AA723802, AA233673,
			nucleotide sequence described by	AI910795, AA527075, AI687053, AI289782,
			the general formula of a-b, where a	AW195947, AA494414, AI680070, AW132045,
			is any integer between 1 to 2506 of	AI368513, AI688692, AW439152, AI688681, C00730,
			SEQ ID NO:2117, b is an integer of	AI697102, AW293340, AA524205, AA514491,
			15 to 2520, where both a and b	AI337294, AI858216, AI857575, AC005837, Y11274
			correspond to the positions of	
	- · - · ·		nucleotide residues shown in SEQ ID	
			NO:2117, and where b is greater	
			than or equal to a + 14.	

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Polynucleotide and Polypeptide Variants

The present invention is directed to variants of the polynucleotide sequence disclosed in SEQ ID NO:X, the complementary strand thereto, and/or the cDNA sequence contained in a deposited clone.

The present invention also encompasses variants of the polypeptide sequence disclosed in SEQ ID NO:Y and/or encoded by a deposited clone.

"Variant" refers to a polynucleotide or polypeptide differing from the polynucleotide or polypeptide of the present invention, but retaining essential properties thereof. Generally, variants are overall closely similar, and, in many regions, identical to the polynucleotide or polypeptide of the present invention.

The present invention is also directed to nucleic acid molecules which comprise, or alternatively consist of, a nucleotide sequence which is at least 80%, 85%, 90%, 95%, 96%, 97%, 98% or 99% identical to, for example, the nucleotide coding sequence in SEQ ID NO:X or the complementary strand thereto, the nucleotide coding sequence contained in a deposited cDNA clone or the complementary strand thereto, a nucleotide sequence encoding the polypeptide of SEQ ID NO:Y, a nucleotide sequence encoding the polypeptide encoded by the cDNA contained in a deposited clone, and/or polynucleotide fragments of any of these nucleic acid molecules (e.g., those fragments described herein). Polynucleotides which hybridize to these nucleic acid molecules under stringent hybridization conditions or alternatively, under lower stringency conditions are also encompassed by the invention, as are polypeptides encoded by these polynucleotides.

The present invention is also directed to polypeptides which comprise, or alternatively consist of, an amino acid sequence which is at least 80%, 85%, 90%, 95%, 96%, 97%, 98%, 99% or 100% identical to, for example, the polypeptide sequence shown in SEQ ID NO:Y, a polypeptide sequence encoded by SEQ ID NO:X or the complement thereof, the polypeptide sequence encoded by the cDNA contained in a deposited clone, and/or polypeptide fragments of any of these polypeptides (e.g., those fragments described herein).

By a nucleic acid having a nucleotide sequence at least, for example, 95% "identical" to a reference nucleotide sequence of the present invention, it is intended that the nucleotide sequence of the nucleic acid is identical to the reference sequence except that the nucleotide sequence may include up to five point mutations per each 100 nucleotides of the reference nucleotide sequence encoding the polypeptide. In other words, to obtain a nucleic acid

having a nucleotide sequence at least 95% identical to a reference nucleotide sequence, up to 5% of the nucleotides in the reference sequence may be deleted or substituted with another nucleotide, or a number of nucleotides up to 5% of the total nucleotides in the reference sequence may be inserted into the reference sequence. The query sequence may be an entire sequence shown in Table 1, the ORF (open reading frame), or any fragment specified as described herein.

As a practical matter, whether any particular nucleic acid molecule or polypeptide is at least 80%, 85%, 90%, 95%, 96%, 97%, 98% or 99% identical to a nucleotide sequence of the presence invention can be determined conventionally using known computer programs. A preferred method for determining the best overall match between a query sequence (a sequence of the present invention) and a subject sequence, also referred to as a global sequence alignment, can be determined using the FASTDB computer program based on the algorithm of Brutlag et al. (Comp. App. Biosci. (1990) 6:237-245). In a sequence alignment the query and subject sequences are both DNA sequences. An RNA sequence can be compared by converting U's to T's. The result of said global sequence alignment is in percent identity. Preferred parameters used in a FASTDB alignment of DNA sequences to calculate percent identity are: Matrix=Unitary, k-tuple=4, Mismatch Penalty=1, Joining Penalty=30, Randomization Group Length=0, Cutoff Score=1, Gap Penalty=5, Gap Size Penalty 0.05, Window Size=500 or the length of the subject nucleotide sequence, whichever is shorter.

If the subject sequence is shorter than the query sequence because of 5' or 3' deletions, not because of internal deletions, a manual correction must be made to the results. This is because the FASTDB program does not account for 5' and 3' truncations of the subject sequence when calculating percent identity. For subject sequences truncated at the 5' or 3' ends, relative to the query sequence, the percent identity is corrected by calculating the number of bases of the query sequence that are 5' and 3' of the subject sequence, which are not matched/aligned, as a percent of the total bases of the query sequence. Whether a nucleotide is matched/aligned is determined by results of the FASTDB sequence alignment. This percentage is then subtracted from the percent identity, calculated by the above FASTDB program using the specified parameters, to arrive at a final percent identity score. This corrected score is what is used for the purposes of the present invention. Only bases outside the 5' and 3' bases of the subject sequence, as displayed by the FASTDB alignment,

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which are not matched/aligned with the query sequence, are calculated for the purposes of manually adjusting the percent identity score.

For example, a 90 base subject sequence is aligned to a 100 base query sequence to determine percent identity. The deletions occur at the 5' end of the subject sequence and therefore, the FASTDB alignment does not show a matched/alignment of the first 10 bases at 5' end. The 10 unpaired bases represent 10% of the sequence (number of bases at the 5' and 3' ends not matched/total number of bases in the query sequence) so 10% is subtracted from the percent identity score calculated by the FASTDB program. If the remaining 90 bases were perfectly matched the final percent identity would be 90%. In another example, a 90 base subject sequence is compared with a 100 base query sequence. This time the deletions are internal deletions so that there are no bases on the 5' or 3' of the subject sequence which are not matched/aligned with the query. In this case the percent identity calculated by FASTDB is not manually corrected. Once again, only bases 5' and 3' of the subject sequence which are not matched/aligned with the query sequence are manually corrected for. No other manual corrections are to made for the purposes of the present invention.

By a polypeptide having an amino acid sequence at least, for example, 95% "identical" to a query amino acid sequence of the present invention, it is intended that the amino acid sequence of the subject polypeptide is identical to the query sequence except that the subject polypeptide sequence may include up to five amino acid alterations per each 100 amino acids of the query amino acid sequence. In other words, to obtain a polypeptide having an amino acid sequence at least 95% identical to a query amino acid sequence, up to 5% of the amino acid residues in the subject sequence may be inserted, deleted, (indels) or substituted with another amino acid. These alterations of the reference sequence may occur at the amino or carboxy terminal positions of the reference amino acid sequence or anywhere between those terminal positions, interspersed either individually among residues in the reference sequence or in one or more contiguous groups within the reference sequence.

As a practical matter, whether any particular polypeptide is at least 80%, 85%, 90%, 95%, 96%, 97%, 98% or 99% identical to, for instance, the amino acid sequences shown in Table 1 or a fragment thereof, or to the amino acid sequence encoded by the cDNA contained in a deposited clone or a fragment thereof, can be determined conventionally using known computer programs. A preferred method for determine the best overall match between a query sequence (a sequence of the present invention) and a subject sequence, also referred to

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as a global sequence alignment, can be determined using the FASTDB computer program based on the algorithm of Brutlag et al. (Comp. App. Biosci.6:237- 245(1990)). In a sequence alignment the query and subject sequences are either both nucleotide sequences or both amino acid sequences. The result of said global sequence alignment is in percent identity. Preferred parameters used in a FASTDB amino acid alignment are: Matrix=PAM 0, k-tuple=2, Mismatch Penalty=1, Joining Penalty=20, Randomization Group Length=0, Cutoff Score=1, Window Size=sequence length, Gap Penalty=5, Gap Size Penalty=0.05, Window Size=500 or the length of the subject amino acid sequence, whichever is shorter.

If the subject sequence is shorter than the query sequence due to N- or C-terminal deletions, not because of internal deletions, a manual correction must be made to the results. This is because the FASTDB program does not account for N- and C-terminal truncations of the subject sequence when calculating global percent identity. For subject sequences truncated at the N- and C-termini, relative to the query sequence, the percent identity is corrected by calculating the number of residues of the query sequence that are N- and Cterminal of the subject sequence, which are not matched/aligned with a corresponding subject residue, as a percent of the total bases of the query sequence. Whether a residue is matched/aligned is determined by results of the FASTDB sequence alignment. percentage is then subtracted from the percent identity, calculated by the above FASTDB program using the specified parameters, to arrive at a final percent identity score. This final percent identity score is what is used for the purposes of the present invention. Only residues to the N- and C-termini of the subject sequence, which are not matched/aligned with the query sequence, are considered for the purposes of manually adjusting the percent identity score. That is, only query residue positions outside the farthest N- and C- terminal residues of the subject sequence.

For example, a 90 amino acid residue subject sequence is aligned with a 100 residue query sequence to determine percent identity. The deletion occurs at the N-terminus of the subject sequence and therefore, the FASTDB alignment does not show a matching/alignment of the first 10 residues at the N-terminus. The 10 unpaired residues represent 10% of the sequence (number of residues at the N- and C- termini not matched/total number of residues in the query sequence) so 10% is subtracted from the percent identity score calculated by the FASTDB program. If the remaining 90 residues were perfectly matched the final percent identity would be 90%. In another example, a 90 residue subject sequence is compared with

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a 100 residue query sequence. This time the deletions are internal deletions so there are no residues at the N- or C-termini of the subject sequence which are not matched/aligned with the query. In this case the percent identity calculated by FASTDB is not manually corrected. Once again, only residue positions outside the N- and C-terminal ends of the subject sequence, as displayed in the FASTDB alignment, which are not matched/aligned with the query sequence are manually corrected for. No other manual corrections are to made for the purposes of the present invention.

The variants may contain alterations in the coding regions, non-coding regions, or both. Especially preferred are polynucleotide variants containing alterations which produce silent substitutions, additions, or deletions, but do not alter the properties or activities of the encoded polypeptide. Nucleotide variants produced by silent substitutions due to the degeneracy of the genetic code are preferred. Moreover, variants in which less than 50, less than 40, less than 30, less than 20, less than 10, or 5-50, 5-25, 5-10, 1-5, or 1-2 amino acids are substituted, deleted, or added in any combination are also preferred. Polynucleotide variants can be produced for a variety of reasons, e.g., to optimize codon expression for a particular host (change codons in the human mRNA to those preferred by a bacterial host such as E. coli).

Naturally occurring variants are called "allelic variants," and refer to one of several alternate forms of a gene occupying a given locus on a chromosome of an organism. (Genes II, Lewin, B., ed., John Wiley & Sons, New York (1985).) These allelic variants can vary at either the polynucleotide and/or polypeptide level and are included in the present invention. Alternatively, non-naturally occurring variants may be produced by mutagenesis techniques or by direct synthesis.

Using known methods of protein engineering and recombinant DNA technology, variants may be generated to improve or alter the characteristics of the polypeptides of the present invention. For instance, one or more amino acids can be deleted from the N-terminus or C-terminus of the colon cancer related polypeptides without substantial loss of biological function. The authors of Ron et al., J. Biol. Chem. 268: 2984-2988 (1993), reported variant KGF proteins having heparin binding activity even after deleting 3, 8, or 27 amino-terminal amino acid residues. Similarly, Interferon gamma exhibited up to ten times higher activity after deleting 8-10 amino acid residues from the carboxy terminus of this protein. (Dobeli et al., J. Biotechnology 7:199-216 (1988).)

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Moreover, ample evidence demonstrates that variants often retain a biological activity similar to that of the naturally occurring protein. For example, Gayle and coworkers (J. Biol. Chem 268:22105-22111 (1993)) conducted extensive mutational analysis of human cytokine IL-1a. They used random mutagenesis to generate over 3,500 individual IL-1a mutants that averaged 2.5 amino acid changes per variant over the entire length of the molecule. Multiple mutations were examined at every possible amino acid position. The investigators found that "[m]ost of the molecule could be altered with little effect on either [binding or biological activity]." (See, Abstract.) In fact, only 23 unique amino acid sequences, out of more than 3,500 nucleotide sequences examined, produced a protein that significantly differed in activity from wild-type.

Furthermore, even if deleting one or more amino acids from the N-terminus or C-terminus of a polypeptide results in modification or loss of one or more biological functions, other biological activities may still be retained. For example, the ability of a deletion variant to induce and/or to bind antibodies which recognize the secreted form will likely be retained when less than the majority of the residues of the secreted form are removed from the N-terminus or C-terminus. Whether a particular polypeptide lacking N- or C-terminal residues of a protein retains such immunogenic activities can readily be determined by routine methods described herein and otherwise known in the art.

Thus, the invention further includes polypeptide variants which show substantial biological activity. Such variants include deletions, insertions, inversions, repeats, and substitutions selected according to general rules known in the art so as have little effect on activity. The present application is directed to nucleic acid molecules at least 80%, 85%, 90%, 95%, 96%, 97%, 98% or 99% or 100% identical to the nucleic acid sequences disclosed herein, (e.g., encoding a polypeptide having the amino acid sequence of an N and/or C terminal deletion), irrespective of whether they encode a polypeptide having functional activity. This is because even where a particular nucleic acid molecule does not encode a polypeptide having functional activity, one of skill in the art would still know how to use the nucleic acid molecule, for instance, as a hybridization probe or a polymerase chain reaction (PCR) primer. Uses of the nucleic acid molecules of the present invention that do not encode a polypeptide having functional activity include, inter alia, (1) isolating a gene or allelic or splice variants thereof in a cDNA library; (2) in situ hybridization (e.g., "FISH") to metaphase chromosomal spreads to provide precise chromosomal location of the gene, as

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described in Verma et al., Human Chromosomes: A Manual of Basic Techniques, Pergamon Press, New York (1988); and (3) Northern Blot analysis for detecting mRNA expression in specific tissues.

Preferred, however, are nucleic acid molecules having sequences at least 80%, 85%, 90%, 95%, 96%, 97%, 98% or 99% or 100% identical to the nucleic acid sequences disclosed herein, which do, in fact, encode a polypeptide having functional activity. By "a polypeptide having functional activity" is intended polypeptides exhibiting activity similar, but not necessarily identical, to a functional activity of the polypeptides of the present invention (e.g., complete (full-length), mature and soluble (e.g., having sequences contained in the extracellular domain) as measured, for example, in a particular immunoassay or biological assay. For example, a functional activity can routinely be measured by determining the ability of a polypeptide of the present invention to bind a ligand. Functional activity may also be measured by determining the ability of a polypeptide, such as cognate ligand which is free or expressed on a cell surface, to induce cells expressing the polypeptide.

Of course, due to the degeneracy of the genetic code, one of ordinary skill in the art will immediately recognize that a large number of the nucleic acid molecules having a sequence at least 80%, 85%, 90%, 95%, 96%, 97%, 98%, or 99%, or 100% identical to, for example, the nucleic acid sequence of the deposited cDNA, the nucleic acid sequence shown in Table 1 (SEQ ID NO:X), or fragments thereof, will encode polypeptides "having functional activity." In fact, since degenerate variants of any of these nucleotide sequences all encode the same polypeptide, in many instances, this will be clear to the skilled artisan even without performing the above described comparison assay. It will be further recognized in the art that, for such nucleic acid molecules that are not degenerate variants, a reasonable number will also encode a polypeptide having functional activity. This is because the skilled artisan is fully aware of amino acid substitutions that are either less likely or not likely to significantly effect protein function (e.g., replacing one aliphatic amino acid with a second aliphatic amino acid), as further described below.

For example, guidance concerning how to make phenotypically silent amino acid substitutions is provided in Bowie et al., "Deciphering the Message in Protein Sequences: Tolerance to Amino Acid Substitutions," Science 247:1306-1310 (1990), wherein the authors indicate that there are two main strategies for studying the tolerance of an amino acid sequence to change.

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The first strategy exploits the tolerance of amino acid substitutions by natural selection during the process of evolution. By comparing amino acid sequences in different species, conserved amino acids can be identified. These conserved amino acids are likely important for protein function. In contrast, the amino acid positions where substitutions have been tolerated by natural selection indicates that these positions are not critical for protein function. Thus, positions tolerating amino acid substitution could be modified while still maintaining biological activity of the protein.

The second strategy uses genetic engineering to introduce amino acid changes at specific positions of a cloned gene to identify regions critical for protein function. For example, site directed mutagenesis or alanine-scanning mutagenesis (introduction of single alanine mutations at every residue in the molecule) can be used. (Cunningham and Wells, Science 244:1081-1085 (1989).) The resulting mutant molecules can then be tested for biological activity.

As the authors state, these two strategies have revealed that proteins are surprisingly tolerant of amino acid substitutions. The authors further indicate which amino acid changes are likely to be permissive at certain amino acid positions in the protein. For example, most buried (within the tertiary structure of the protein) amino acid residues require nonpolar side chains, whereas few features of surface side chains are generally conserved. Moreover, tolerated conservative amino acid substitutions involve replacement of the aliphatic or hydrophobic amino acids Ala, Val, Leu and Ile; replacement of the hydroxyl residues Ser and Thr; replacement of the acidic residues Asp and Glu; replacement of the amide residues Asn and Gln, replacement of the basic residues Lys, Arg, and His; replacement of the aromatic residues Phe, Tyr, and Trp, and replacement of the small-sized amino acids Ala, Ser, Thr, Met, and Gly. Besides conservative amino acid substitution, variants of the present invention include (i) substitutions with one or more of the non-conserved amino acid residues, where the substituted amino acid residues may or may not be one encoded by the genetic code, or (ii) substitution with one or more of amino acid residues having a substituent group, or (iii) fusion of the mature polypeptide with another compound, such as a compound to increase the stability and/or solubility of the polypeptide (for example, polyethylene glycol), or (iv) fusion of the polypeptide with additional amino acids, such as, for example, an IgG Fc fusion region peptide, or leader or secretory sequence, or a sequence facilitating purification. Such variant

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polypeptides are deemed to be within the scope of those skilled in the art from the teachings herein.

For example, polypeptide variants containing amino acid substitutions of charged amino acids with other charged or neutral amino acids may produce proteins with improved characteristics, such as less aggregation. Aggregation of pharmaceutical formulations both reduces activity and increases clearance due to the aggregate's immunogenic activity. (Pinckard et al., Clin. Exp. Immunol. 2:331-340 (1967); Robbins et al., Diabetes 36: 838-845 (1987); Cleland et al., Crit. Rev. Therapeutic Drug Carrier Systems 10:307-377 (1993).)

A further embodiment of the invention relates to a polypeptide which comprises the amino acid sequence of a polypeptide having an amino acid sequence which contains at least one amino acid substitution, but not more than 50 amino acid substitutions, even more preferably, not more than 40 amino acid substitutions, still more preferably, not more than 30 amino acid substitutions, and still even more preferably, not more than 20 amino acid substitutions. Of course, in order of ever-increasing preference, it is highly preferable for a polypeptide to have an amino acid sequence which comprises the amino acid sequence of a polypeptide of SEQ ID NO:Y, in order of ever-increasing preference, which contains at least one, but not more than 10, 9, 8, 7, 6, 5, 4, 3, 2 or 1 amino acid substitutions. In specific embodiments, the number of additions, substitutions, and/or deletions in the amino acid sequence of SEQ ID NO:Y or fragments thereof (e.g., the mature form and/or other fragments described herein), and/or the amino acid sequence encoded by the deposited clone or fragments thereof, is 1-5, 5-10, 5-25, 5-50, 10-50 or 50-150, conservative amino acid substitutions are preferable.

Polynucleotide and Polypeptide Fragments

The present invention is also directed to polynucleotide fragments of the polynucleotides of the invention. In the present invention, a "polynucleotide fragment" refers to a short polynucleotide having a nucleic acid sequence which: is a portion of the cDNA contained in a depostied cDNA clone; or is a portion of a polynucleotide sequence encoding the polypeptide encoded by the cDNA contained in a deposited cDNA clone; or is a portion of the polynucleotide sequence in SEQ ID NO:X or the complementary strand thereto; or is a polynucleotide sequence encoding a portion of the polypeptide of SEQ ID NO:Y; or is a polynucleotide sequence encoding a portion of a polypeptide encoded by SEQ ID NO:X or

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the complementary strand thereto. The nucleotide fragments of the invention are preferably at least about 15 nt, and more preferably at least about 20 nt, still more preferably at least about 30 nt, and even more preferably, at least about 40nt, at least about 50 nt, at least about 75 nt, or at least about 150 nt in length. A fragment "at least 20 nt in length," for example, is intended to include 20 or more contiguous bases from the cDNA sequence contained in a deposited clone or the nucleotide sequence shown in SEQ ID NO:X or the complementary strand thereto. In this context "about" includes the particularly recited value, a value larger or smaller by several (5, 4, 3, 2, or 1) nucleotides, at either terminus or at both termini. These nucleotide fragments have uses that include, but are not limited to, as diagnostic probes and primers as discussed herein. Of course, larger fragments (e.g., at least 50, 150, 200, 250, 500, 600, 1000 or 2000 nucleotides in length) are also encompassed by the invention.

Moreover, representative examples of polynucleotide fragments of the invention, include, for example, fragments comprising, or alternatively consisting of, a sequence from about nucleotide number 1-50, 51-100, 101-150, 151-200, 201-250, 251-300, 301-350, 351-400, 401-450, 451-500, 501-550, 551-600, 651-700, 701-750, 751-800, 800-850, 851-900, 901-950, 951-1000, 1001-1050, 1051-1100, 1101-1150, 1151-1200, 1201-1250, 1251-1300, 1301-1350, 1351-1400, 1401-1450, 1451-1500, 1501-1550, 1551-1600, 1601-1650, 1651-1700, 1701-1750, 1751-1800, 1801-1850, 1851-1900, 1901-1950, 1951-2000, 2001-2050, 2051-2100, 2101-2150, 2151-2200, 2201-2250, 2251-2300, 2301-2350, 2351-2400, 2401-2450, 2451-2500, 2501-2550, 2551-2600, 2651-2700, 2701-2750, 2751-2800, 2800-2850, 2851-2900, 2901-2950, 2951-3000, 3001-3050, 3051-3100 and 3101 to the end of SEQ ID NO:X, or the complementary strand thereto, or the cDNA contained in the deposited clone. In this context "about" includes the particularly recited ranges, and ranges larger or smaller by several (5, 4, 3, 2, or 1) nucleotides, at either terminus or at both termini. Preferably, these fragments encode a polypeptide which has biological activity. More preferably, these polynucleotides can be used as probes or primers as discussed herein. Polynucleotides which hybridize to these nucleic acid molecules under stringent hybridization conditions or lower stringency conditions are also encompassed by the invention, as are polypeptides encoded by these polynucleotides.

Moreover, representative examples of polynucleotide fragments of the invention, include, for example, fragments comprising, or alternatively consisting of, a sequence from about nucleotide number 1-50, 51-100, 101-150, 151-200, 201-250, 251-300, 301-350, 351-

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400, 401-450, 451-500, 501-550, 551-600, 651-700,701-750, 751-800, 800-850, 851-900, 901-950, 951-1000, 1001-1050, 1051-1100, 1101-1150, 1151-1200, 1201-1250, 1251-1300, 1301-1350, 1351-1400, 1401-1450, 1451-1500, 1501-1550, 1551-1600, 1601-1650, 1651-1700, 1701-1750, 1751-1800, 1801-1850, 1851-1900, 1901-1950, 1951-2000, 2001-2050, 2051-2100, 2101-2150, 2151-2200, 2201-2250, 2251-2300, 2301-2350, 2351-2400, 2401-2450, 2451-2500, 2501-2550, 2551-2600, 2601-2650, 2651-2700, 2701-2750, 2751-2800, 2801-2850, 2851-2900, 2901-2950, 2951-3000, 3001-3050, 3051-3100 and 3101 to the end of the cDNA nucleotide sequence contained in the deposited cDNA clone, or the complementary strand thereto. In this context "about" includes the particularly recited range, or a range larger or smaller by several (5, 4, 3, 2, or 1) nucleotides, at either terminus or at both termini. Preferably, these fragments encode a polypeptide which has a functional activity (e.g., biological activity) of the polypeptide encoded by the cDNA nucleotide sequence contained in the deposited cDNA clone. More preferably, these fragments can be used as probes or primers as discussed herein. Polynucleotides which hybridize to one or more of these fragments under stringent hybridization conditions or alternatively, under lower stringency conditions, are also encompassed by the invention, as are polypeptides encoded by these polynucleotides or fragments.

In the present invention, a "polypeptide fragment" refers to an amino acid sequence which is a portion of that contained in SEQ ID NO:Y, encoded by SEQ ID NO:X or the complement thereof and/or encoded by the cDNA contained in the deposited clone. Protein (polypeptide) fragments may be "free-standing," or comprised within a larger polypeptide of which the fragment forms a part or region, most preferably as a single continuous region. Representative examples of polypeptide fragments of the invention, include, for example, fragments comprising, or alternatively consisting of, from about amino acid number 1-20, 21-40, 41-60, 61-80, 81-100, 102-120, 121-140, 141-160, 161-180, 181-200, 201-220, 221-240, 241-260, 261-280, 281-300, 301-320, 321-340, 341-360, 361-380, 381-400, 401-420, 421-440, 441-460, 461-480, 481-500, 501-520, 521-540, 541-560, 561-580, 581-600, 601-620, 621-640, 641-660, 661-680, 681-700, 701-720, 721-740, 741-760, 761-780, 781-800, 801-820, 821-840, 841-860 and 861 to the end of SEQ ID NO:Y. Moreover, polypeptide fragments can be about 10, 15, 20, 25, 30, 35, 40, 45, 50, 55, 60, 65, 70, 75, 80, 85, 90, 100, 110, 120, 130, 140, or 150 amino acids in length. In this context "about" includes the particularly recited ranges or values, and ranges or values larger or smaller by several (5, 4, 3,

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2, or 1) amino acids, at either extreme or at both extremes. Polynucleotides encoding these polypeptides are also encompassed by the invention.

Even if deletion of one or more amino acids from the N-terminus of a protein results in modification of loss of one or more biological functions of the protein, other functional activities (e.g., biological activities, ability to multimerize, ability to bind a ligand) may still be retained. For example, the ability of shortened muteins to induce and/or bind to antibodies which recognize the complete or mature forms of the polypeptides generally will be retained when less than the majority of the residues of the complete or mature polypeptide are removed from the N-terminus. Whether a particular polypeptide lacking N-terminal residues of a complete polypeptide retains such immunologic activities can readily be determined by routine methods described herein and otherwise known in the art. It is not unlikely that a mutein with a large number of deleted N-terminal amino acid residues may retain some biological or immunogenic activities. In fact, peptides composed of as few as six amino acid residues may often evoke an immune response.

Accordingly, polypeptide fragments include the secreted protein as well as the mature form. Further preferred polypeptide fragments include the secreted protein or the mature form having a continuous series of deleted residues from the amino or the carboxy terminus, or both. For example, any number of amino acids, ranging from 1-60, can be deleted from the amino terminusof either the secreted polypeptide or the mature form. Similarly, any number of amino acids, ranging from 1-30, can be deleted from the carboxy terminus of the secreted protein or mature form. Furthermore, any combination of the above amino and carboxy terminus deletions are preferred. Similarly,polynucleotides encoding these polypeptide fragments are also preferred.

The present invention further provides polypeptides having one or more residues deleted from the amino terminus of the amino acid sequence of a polypeptide disclosed herein (e.g., a polypeptide of SEQ ID NO:Y, and/or a polypeptide encoded by the cDNA contained in a deposited clone). In particular, N-terminal deletions may be described by the general formula m-q, where q is a whole integer representing the total number of amino acid residues in a polypeptide of the invention (e.g., the polypeptide disclosed in SEQ ID NO:Y), and m is defined as any integer ranging from 2 to q-6. Polynucleotides encoding these polypeptides are also encompassed by the invention.

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Also as mentioned above, even if deletion of one or more amino acids from the C-terminus of a protein results in modification of loss of one or more biological functions of the protein, other functional activities (e.g., biological activities, ability to multimerize, ability to bind a ligand) may still be retained. For example the ability of the shortened mutein to induce and/or bind to antibodies which recognize the complete or mature forms of the polypeptide generally will be retained when less than the majority of the residues of the complete or mature polypeptide are removed from the C-terminus. Whether a particular polypeptide lacking C-terminal residues of a complete polypeptide retains such immunologic activities can readily be determined by routine methods described herein and otherwise known in the art. It is not unlikely that a mutein with a large number of deleted C-terminal amino acid residues may retain some biological or immunogenic activities. In fact, peptides composed of as few as six amino acid residues may often evoke an immune response.

Accordingly, the present invention further provides polypeptides having one or more residues from the carboxy terminus of the amino acid sequence of a polypeptide disclosed herein (e.g., a polypeptide of SEQ ID NO:Y, a polypeptide encoded by the polynucleotide sequence contained in SEQ ID NO:X, and/or a polypeptide encoded by the cDNA contained in deposited cDNA clone referenced in Table 1). In particular, C-terminal deletions may be described by the general formula 1-n, where n is any whole integer ranging from 6 to q-1, and where n corresponds to the position of an amino acid residue in a polypeptide of the invention. Polynucleotides encoding these polypeptides are also encompassed by the invention.

In addition, any of the above described N- or C-terminal deletions can be combined to produce a N- and C-terminal deleted polypeptide. The invention also provides polypeptides having one or more amino acids deleted from both the amino and the carboxyl termini, which may be described generally as having residues m-n of a polypeptide encoded by SEQ ID NO:X (e.g., including, but not limited to the preferred polypeptide disclosed as SEQ ID NO:Y), or the cDNA contained in a deposited clone, and/or the complement thereof, where n and m are integers as described above. Polynucleotides encoding these polypeptides are also encompassed by the invention.

Any polypeptide sequence contained in the polypeptide of SEQ ID NO:Y, encoded by the polynucleotide sequences set forth as SEQ ID NO:X or the complement thereof, or encoded by the cDNA in the related cDNA clone contained in the deposit may be analyzed to

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determine certain preferred regions of the polypeptide. For example, the amino acid sequence of a polypeptide encoded by a polynucleotide sequence of SEQ ID NO:X or the complement thereof, or the cDNA in a deposited cDNA clone may be analyzed using the default parameters of the DNASTAR computer algorithm (DNASTAR, Inc., 1228 S. Park St., Madison, WI 53715 USA; http://www.dnastar.com/).

Polypeptide regions that may be routinely obtained using the DNASTAR computer algorithm include, but are not limited to, Garnier-Robson alpha-regions, beta-regions, turn-regions, and coil-regions, Chou-Fasman alpha-regions, beta-regions, and turn-regions, Kyte-Doolittle hydrophilic regions and hydrophobic regions, Eisenberg alpha- and beta-amphipathic regions, Karplus-Schulz flexible regions, Emini surface-forming regions and Jameson-Wolf regions of high antigenic index. Among highly preferred polynucleotides of the invention in this regard are those that encode polypeptides comprising regions that combine several structural features, such as several (e.g., 1, 2, 3 or 4) of the features set out above.

Additionally, Kyte-Doolittle hydrophilic regions and hydrophobic regions, Emini surface-forming regions, and Jameson-Wolf regions of high antigenic index (i.e., containing four or more contiguous amino acids having an antigenic index of greater than or equal to 1.5, as identified using the default parameters of the Jameson-Wolf program) can routinely be used to determine polypeptide regions that exhibit a high degree of potential for antigenicity. Regions of high antigenicity are determined from data by DNASTAR analysis by choosing values which represent regions of the polypeptide which are likely to be exposed on the surface of the polypeptide in an environment in which antigen recognition may occur in the process of initiation of an immune response.

Preferred polypeptide fragments of the invention are fragments comprising, or alternatively consisting of, an amino acid sequence that displays a functional activity of the polypeptide sequence of which the amino acid sequence is a fragment.

By a polypeptide demonstrating a "functional activity" is meant, a polypeptide capable of displaying one or more known functional activities associated with a full-length (complete) protein of the invention. Such functional activities include, but are not limited to, biological activity, antigenicity [ability to bind (or compete with a polypeptide for binding) to an anti-polypeptide antibody], immunogenicity (ability to generate antibody which binds to

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a specific polypeptide of the invention), ability to form multimers with polypeptides of the invention, and ability to bind to a receptor or ligand for a polypeptide.

Other preferred polypeptide fragments are biologically active fragments. Biologically active fragments are those exhibiting activity similar, but not necessarily identical, to an activity of the polypeptide of the present invention. The biological activity of the fragments may include an improved desired activity, or a decreased undesirable activity.

In preferred embodiments, polypeptides of the invention comprise, or alternatively consist of, one, two, three, four, five or more of the antigenic fragments of the polypeptide of SEQ ID NO:Y, or portions thereof. Polynucleotides encoding these polypeptides are also encompassed by the invention.

Table 8

Contig ID/	Epitopes
Sequence	
ID	
390631	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4278 as residues: Asn-1 to Asn-6.
410299	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4279 as residues: Trp-26 to Met-31.
456200	Preferred epitopes include those comprising a sequence shown in SEQ
15155	ID NO. 4280 as residues: Pro-16 to His-26, Arg-45 to Gly-51.
471563	Preferred epitopes include those comprising a sequence shown in SEQ
400121	ID NO. 4283 as residues: Gly-37 to Glu-47.
488131	Preferred epitopes include those comprising a sequence shown in SEQ
500606	ID NO. 4284 as residues: Met-26 to Leu-32, Gly-41 to Asn-46.
500696	Preferred epitopes include those comprising a sequence shown in SEQ
506406	ID NO. 4286 as residues: Lys-16 to Glu-31, Ser-47 to Glu-54.
506406	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4288 as residues: Thr-110 to Tyr-118.
506619	Preferred epitopes include those comprising a sequence shown in SEQ
300019	ID NO. 4289 as residues: Cys-50 to Phe-57, Phe-69 to Asp-76, Ser-89
	to Gln-104, Glu-145 to Leu-153.
507852	Preferred epitopes include those comprising a sequence shown in SEQ
50,052	ID NO. 4290 as residues: Glu-8 to Trp-18, Arg-46 to Ala-51.
509423	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4291 as residues: Tyr-50 to Ser-56, His-58 to Tyr-65.
524721	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4294 as residues: Pro-1 to Ser-8.
524901	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4295 as residues: Leu-34 to Lys-39, Lys-57 to Gly-63.
527600	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4296 as residues: Val-28 to Gly-34, His-57 to His-63.
529050	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4298 as residues: Asn-2 to Lys-8.
529465	Preferred epitopes include those comprising a sequence shown in SEQ
622010	ID NO. 4299 as residues: Ala-12 to Gln-24.
532810	Preferred epitopes include those comprising a sequence shown in SEQ
541126	ID NO. 4302 as residues: Pro-1 to Trp-7, Glu-124 to Trp-130.
541126	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4304 as residues: Thr-1 to Asn-10, Ala-72 to Gly-77, Val-84 to
	Gly-90.
542268	Preferred epitopes include those comprising a sequence shown in SEQ
3-2200	ID NO. 4305 as residues: Pro-34 to Pro-40, Pro-45 to Ser-50, Gly-73 to
	Gly-82.
547920	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4306 as residues: Pro-28 to Thr-35.
552465	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4310 as residues: Pro-4 to Gly-10, Thr-17 to Leu-29, Pro-53 to
	Gly-58, Gln-78 to Lys-86, Pro-88 to Lys-94, His-137 to Gly-142.

554260	DC1
554369	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4311 as residues: Gln-20 to Gln-27.
557152	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4312 as residues: Ser-69 to Pro-74.
557230	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4313 as residues: Pro-21 to Cys-31, Val-34 to Gly-42.
570796	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4315 as residues: Glu-34 to Ala-39.
573181	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4316 as residues: Gly-4 to Arg-11, Gly-17 to Ala-24.
573793	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4318 as residues: Glu-4 to Ser-9.
573796	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4319 as residues: Pro-4 to Asn-13, Asn-57 to Arg-66, Pro-89 to Asn-99.
574927	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4321 as residues: Asp-1 to Ile-6, Pro-37 to Gln-42, Pro-61 to Trp-68.
575139	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4322 as residues: Met-2 to Asp-9.
575591	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4323 as residues: Ala-2 to Gly-11.
577390	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4325 as residues: Glu-53 to Leu-58, Gln-60 to Glu-65.
577685	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4326 as residues: Ile-5 to Gln-12, Leu-42 to Asn-51.
578660	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4328 as residues: His-1 to Phe-6, Val-11 to Arg-23.
580860	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4329 as residues: Ser-14 to Asn-22.
581143	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4330 as residues: Ile-1 to Gly-6.
584899	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4331 as residues: Ala-29 to Asn-35.
600669	Preferred epitopes include those comprising a sequence shown in SEO
611839	ID NO. 4332 as residues: Cys-1 to Ala-18, Cys-55 to Ile-61. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4333 as residues: Arg 35 to Chr. 41
614078	ID NO. 4333 as residues: Arg-35 to Gly-41. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4334 as residues: Glu-8 to Leu-14.
630230	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4338 as residues: Arg-77 to Lys-83.
637605	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4340 as residues: Ser-1 to Val-11.
638125	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4341 as residues: Ser-7 to Glu-12, Pro-20 to Ser-26, Arg-31 to Glu-43, Ala-69 to Glu-80, Val-90 to His-95, Pro-100 to Ser-107, Ser-109 to Glu-115, Ala-117 to Arg-124.
638249	Preferred epitopes include those comprising a sequence shown in SEQ

	ID NO. 4343 as residues: Asp-1 to Pro-28, Gln-73 to Ser-79, Ile-91 to Gly-96, Tyr-99 to Asp-109, Gln-183 to Pro-193, Val-249 to Thr-261.
638319	Preferred epitopes include those comprising a sequence shown in SEQ
038319	ID NO. 4344 as residues: Gly-23 to Gly-28, Asp-35 to Gln-53.
651380	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4345 as residues: Thr-16 to Lys-35, Lys-46 to Arg-51.
651876	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4346 as residues: Arg-1 to Asp-12, Pro-25 to Ala-34, Ala-50 to
	Gly-55, Glu-66 to Lys-86.
653175	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4347 as residues: Thr-45 to Asn-50.
655544	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4348 as residues: Arg-2 to Asp-18, Leu-45 to Leu-51.
656722	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4349 as residues: Gln-21 to Leu-38.
659801	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4350 as residues: Gly-2 to Gly-20, Pro-45 to Ala-51, Glu-105 to
	Gln-112, Gln-117 to Glu-122, Ala-207 to Leu-215.
660020	Preferred epitopes include those comprising a sequence shown in SEQ
661401	ID NO. 4351 as residues: Ser-40 to Thr-52.
664481	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4353 as residues: Gly-1 to Glu-15, Phe-20 to Tyr-25, Phe-53 to
665154	Asn-58, Glu-82 to Lys-93.
665154	Preferred epitopes include those comprising a sequence shown in SEQ
((8040	ID NO. 4354 as residues: Pro-18 to Arg-29.
668040	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4356 as residues: Glu-5 to Ala-14, Arg-69 to Ala-76, Ala-114 to Glu-120, Ser-132 to Leu-137.
668717	Preferred epitopes include those comprising a sequence shown in SEQ
000/1/	ID NO. 4358 as residues: Arg-3 to Gly-12, Ala-51 to Asp-65, Leu-78 to
	Glu-84, Arg-118 to Asp-131, Leu-157 to Asp-168.
671361	Preferred epitopes include those comprising a sequence shown in SEQ
0/1501	ID NO. 4360 as residues: Asn-1 to Ser-6, Glu-15 to Gln-20.
674203	Preferred epitopes include those comprising a sequence shown in SEQ
07.1203	ID NO. 4361 as residues: Gly-7 to Ile-13.
674745	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4362 as residues: Val-17 to Arg-26, Lys-38 to Leu-48, Gln-129
	to Trp-136, Gln-258 to Leu-263, Ala-272 to Glu-284, Pro-380 to Asp-
	391.
674761	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4363 as residues: Ala-14 to His-19.
677212	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4364 as residues: Gly-1 to Ser-14, Asn-29 to Trp-34, Lys-50 to
	Arg-60.
685895	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4366 as residues: Arg-28 to Ser-33.
688040	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4367 as residues: Thr-2 to Ser-7, Pro-132 to Asp-138, Ile-161 to
	Pro-170, Pro-212 to Asn-217, Gly-280 to Gln-313, Ser-332 to His-337,

	Apr. 266 to Ch. 272
(00044	Asn-366 to Gly-372.
688044	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4368 as residues: Asn-33 to Pro-55, Lys-67 to Arg-74, Gly-85
	to Tyr-94, Arg-101 to Pro-115, Ser-123 to Cys-129, Pro-155 to Val-162,
- (2112	Pro-172 to Cys-184.
691124	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4370 as residues: Pro-27 to Arg-35.
691721	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4371 as residues: Lys-23 to Gln-29, Gly-59 to Asn-77.
693582	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4372 as residues: Lys-12 to Lys-17.
696007	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4373 as residues: Gln-93 to Arg-101, Tyr-104 to Thr-113. His-
	134 to Gln-145, Ser-154 to Gln-165, Val-231 to Pro-248.
703700	Preferred epitopes include those comprising a sequence shown in SEO
	ID NO. 4377 as residues: Lys-1 to Ser-21.
705461	Preferred epitopes include those comprising a sequence shown in SEO
	ID NO. 4378 as residues: Ala-53 to Glu-59, Thr-69 to Gln-77, Glu-107
	to Trp-114.
707464	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4382 as residues: Glu-1 to Tyr-14, Lys-41 to Arg-51, Thr-54 to
	Arg-73, Gly-77 to Thr-84, Thr-92 to Ser-100, Gln-107 to Arg-112, Ala-
	114 to Ser-141.
709015	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4383 as residues: Pro-62 to Ser-67.
711878	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4387 as residues: Ser-3 to Lys-10.
712638	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4388 as residues: Leu-31 to His-36, Val-94 to Phe-105.
715343	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4392 as residues: Phe-7 to Ile-12, Leu-17 to Ser-24.
716212	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4393 as residues: Ser-1 to Trp-6, Pro-8 to Pro-21, Arg-60 to
	Asp-65, Tyr-70 to Lys-80, Lys-116 to Met-121.
717222	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4394 as residues: Glu-40 to Ala-45, Pro-66 to Ser-80, Gly-99 to
	Ala-107.
719829	Preferred epitopes include those comprising a sequence shown in SEQ
1	ID NO. 4396 as residues: Leu-15 to Cys-20.
721985	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4397 as residues: Asp-1 to Leu-19.
722249	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4398 as residues: Ala-54 to Gly-59, Ser-67 to Gly-78, Ala-131
	to Pro-136, Pro-151 to His-157, Pro-172 to Asn-181, His-183 to Gln-
	192, Ala-200 to Asn-208, Thr-220 to Ile-226, Glu-335 to Arg-341, Ser-
	397 to Cys-404, Lys-415 to Phe-423, Lys-432 to Leu-437.
722258	Preferred epitopes include those comprising a sequence shown in SEQ
, 22250	ID NO. 4399 as residues: Trp-15 to Ala-24, Arg-38 to Glu-45, Tyr-51
	to Gly-59.
	10 013 37.

ID NO. 4401 as residues: Leu-23 to Asn-32. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4402 as residues: Asn-1 to Ser-9, Leu-49 to Leu-64, Leu-68 to Arg-73, Lys-83 to Thr-90. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4404 as residues: Val-36 to Lys-51, Asn-59 to Asn-76, Val-91 to Lys-107, Leu-112 to Cys-135, Arg-140 to Lys-150, Pro-157 to Glu-173, Thr-188 to Lys-201, Lys-207 to Ile-226, Leu-234 to Thr-258, Glu-260 to Ile-268, Ser-275 to Lys-286, Val-288 to Glu-299. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4407 as residues: Lys-8 to His-18. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4411 as residues: Lys-34 to Ser-39. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4412 as residues: His-1 to Gln-6, Glu-19 to Val-26. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4413 as residues: Asn-7 to Lys-13. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4414 as residues: Pro-10 to Trp-18. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4416 as residues: Pro-10 to Trp-18. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4416 as residues: Pro-10 to Trp-18. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4416 as residues: Pro-10 to Trp-18. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4416 as residues: Pro-11 to Trp-13. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4420 as residues: Asp-10 to Sp-118, Trp-17, Gly-134 to Lys-14. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4420 as residues: Asp-10 to Lys-11. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4422 as residues: Asp-10 to Asp-28, Ser-34 to Asp-40. Preferred epitopes include those comprising a sequence show	725110	Preferred epitopes include those comprising a sequence shown in SEQ
ID NO. 4402 as residues: Asn-1 to Ser-9, Leu-49 to Leu-64, Leu-68 to Arg-73, Lys-83 to Thr-90. 727365 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4404 as residues: Val-36 to Lys-51, Asn-59 to Asn-76, Val-91 to Lys-107, Leu-112 to Cys-135, Arg-140 to Lys-150, Pro-157 to Glu-173, Thr-188 to Lys-201, Lys-220 to Ile-226, Leu-234 to Thr-258, Glu-260 to Ile-268, Ser-275 to Lys-286, Val-288 to Glu-299. 731881 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4407 as residues: Lys-8 to His-18. 734012 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4411 as residues: Lys-8 to His-18. 735603 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4412 as residues: Lys-34 to Ser-39. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4413 as residues: Asn-7 to Lys-13. 741134 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4414 as residues: Pro-10 to Trp-18. 741804 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4416 as residues: Pro-10 to Trp-18. 742220 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4416 as residues: Asp-21 to Ser-30, His-37 to Lys-48, Phe-75 to Arg-82. 742220 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4418 as residues: Asp-1 to Lys-11. 745368 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4420 as residues: Asp-1 to Lys-14. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4423 as residues: Asp-1 to Lys-34 to Asp-37. 750486 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4423 as residues: Shy-1 to Lys-28, Ser-34 to Asp-40. 75119 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4423 as residues: Shy-1 to Ayp-28, Ser-34 to Asp-40. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4423 as residues: Shy-1 to Alg-28, Ser-34		ID NO. 4401 as residues: Leu-23 to Asn-32.
Arg-73, Lys-83 to Thr-90. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4404 as residues: Val-36 to Lys-51, Asn-59 to Asn-76, Val-91 to Lys-107, Leu-112 to Cys-135, Arg-140 to Lys-150, Pro-157 to Glu-173, Thr-188 to Lys-201, Lys-207 to Ile-226, Leu-234 to Thr-258, Glu-260 to Ile-268, Ser-275 to Lys-286, Val-288 to Glu-299. 731881 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4407 as residues: Lys-8 to His-18. 734012 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4411 as residues: Lys-34 to Ser-39. 735603 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4412 as residues: His-1 to Glin-6, Gliu-19 to Val-26. 739061 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4413 as residues: Asn-7 to Lys-13. 741134 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4414 as residues: Pro-10 to Trp-18. 741804 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4416 as residues: Asn-21 to Ser-30, His-37 to Lys-48, Phe-75 to Arg-82. 742220 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4417 as residues: Val-17 to Pro-23, Ser-72 to His-79, Thr-93 to Ile-100, Pro-102 to Asp-108, Asn-111 to Tyr-117, Gly-134 to Lys-141. 744605 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4420 as residues: Asp-1 to Lys-11. 750486 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4420 as residues: Lys-10 to Ser-16, Pro-30 to Arg-37. 750486 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4423 as residues: Asp-21 to Asp-28, Ser-34 to Asp-40. 751119 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4423 as residues: Gly-1 to Gly-13, Gly-18 to Glu-29. 753226 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4423 as residues: Gly-1 to Arg-9, Asn-51 to Cys-57, Cys-125 to Leu-137, Cys-153 to Trp-166, Leu-18	725201	
727365 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4404 as residues: Val-36 to Lys-51, Asn-59 to Asn-76, Val-91 to Lys-107, Leu-112 to Cys-135, Arg-140 to Lys-150, Pro-157 to Glu-173, Thr-188 to Lys-201, Lys-207 to Ile-226, Leu-234 to Thr-258, Glu-260 to Ile-268, Ser-275 to Lys-286, Val-288 to Glu-299. 731881 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4407 as residues: Lys-8 to His-18. 734012 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4411 as residues: Lys-34 to Ser-39. 735603 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4412 as residues: Lys-34 to Ser-39. 739061 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4413 as residues: Asn-7 to Lys-13. 741134 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4414 as residues: Asn-7 to Lys-13. 741804 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4416 as residues: Pro-10 to Trp-18. 742220 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4416 as residues: Asp-21 to Ser-30, His-37 to Lys-48, Phe-75 to Arg-82. 742220 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4416 as residues: Val-17 to Pro-23, Ser-72 to His-79, Thr-93 to IIe-100, Pro-102 to Asp-108, Asn-111 to Tyr-117, Gly-134 to Lys-141. 744605 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4418 as residues: Val-10 to Ser-16, Pro-30 to Arg-37. 750486 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4422 as residues: Lys-10 to Ser-16, Pro-30 to Arg-37. 750486 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4422 as residues: Ser-10 to Asp-28, Ser-34 to Asp-40. 75119 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4423 as residues: Asp-1 to Arg-9, Asn-51 to Cys-57, Cys-125 to Leu-137, Cys-153 to Trp-166, Leu-181 to Glu-186, Ser-207 to Thr-212. 756		
ID NO. 4404 as residues: Val-36 to Lys-51, Asn-59 to Asn-76, Val-91 to Lys-107, Leu-112 to Cys-135, Arg-140 to Lys-150, Pro-157 to Glu-173, Th-188 to Lys-201, Lys-207 to Ile-226, Leu-234 to Thr-258, Glu-260 to Ile-268, Ser-275 to Lys-286, Val-288 to Glu-299. 731881 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4407 as residues: Lys-8 to His-18. 734012 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4411 as residues: Lys-34 to Ser-39. 735603 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4412 as residues: His-1 to Gln-6, Glu-19 to Val-26. 739061 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4413 as residues: Asn-7 to Lys-13. 741134 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4414 as residues: Pro-10 to Trp-18. 741804 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4416 as residues: Asn-7 to Lys-43, His-37 to Lys-48, Phe-75 to Arg-82. 742220 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4416 as residues: Val-17 to Pro-23, Ser-72 to His-79, Thr-93 to Ile-100, Pro-102 to Asp-108, Asn-111 to Tyr-117, Gly-134 to Lys-141. 744605 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4420 as residues: Asp-1 to Lys-11. 750486 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4420 as residues: Asp-1 to Asp-13, Ser-34 to Asp-40. 751119 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4423 as residues: Asp-1 to Asp-28, Ser-34 to Asp-40. 751226 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4423 as residues: Asp-1 to Asp-28, Ser-34 to Asp-40. 751226 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4423 as residues: Gly-1 to Gly-13, Gly-18 to Gly-29. 75226 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4425 as residues: Gly-1 to Asp-24. 75227 Preferred epitopes inc		
to Lys-107, Leu-112 to Cys-135, Arg-140 to Lys-150, Pro-157 to Glu- 173, Thr-188 to Lys-201, Lys-207 to Ile-226, Leu-234 to Thr-258, Glu- 260 to Ile-268, Ser-275 to Lys-286, Val-288 to Glu-299. 731881 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4407 as residues: Lys-8 to His-18. 734012 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4411 as residues: Lys-34 to Ser-39. 735603 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4412 as residues: His-1 to Gln-6, Glu-19 to Val-26. 739061 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4413 as residues: Asn-7 to Lys-13. 741134 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4414 as residues: Pro-10 to Trp-18. 741804 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4414 as residues: Asp-21 to Ser-30, His-37 to Lys-48, Phe-75 to Arg-82. 742220 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4417 as residues: Val-17 to Pro-23, Ser-72 to His-79, Thr-93 to Inle-100, Pro-102 to Asp-108, Asn-111 to Tyr-117, Gly-134 to Lys-141. 744605 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4420 as residues: Asp-1 to Lys-11. 745368 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4420 as residues: Lys-10 to Ser-16, Pro-30 to Arg-37. 750486 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4423 as residues: Asp-21 to Asp-28, Ser-34 to Asp-40. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4425 as residues: Gly-1 to Gly-13, Gly-18 to Glu-29, 753226 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4425 as residues: Ser-1 to Arg-9, Asn-51 to Cys-57, Cys-125 to Leu-137, Cys-153 to Trp-166, Leu-181 to Glu-186, Ser-207 to Thr-212. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4427 as residues: Gly-1 to His-10, His-21 to Asp-32. Prefe	727365	Preferred epitopes include those comprising a sequence shown in SEQ
173, Thr-188 to Lys-201, Lys-207 to Ile-226, Leu-234 to Thr-258, Glu-260 to Ile-268, Ser-275 to Lys-286, Val-288 to Glu-299. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4407 as residues: Lys-8 to His-18. 734012 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4411 as residues: Lys-34 to Ser-39. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4412 as residues: His-1 to Gln-6, Glu-19 to Val-26. 739061 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4413 as residues: Asn-7 to Lys-13. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4414 as residues: Pro-10 to Trp-18. 741804 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4416 as residues: Asn-21 to Ser-30, His-37 to Lys-48, Phe-75 to Arg-82. 742220 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4417 as residues: Val-17 to Pro-23, Ser-72 to His-79, Thr-93 to Ile-100, Pro-102 to Asp-108, Asn-111 to Tyr-117, Gly-134 to Lys-141. 744605 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4418 as residues: Asp-1 to Lys-11. 745368 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4420 as residues: Lys-10 to Ser-16, Pro-30 to Arg-37. 750486 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4422 as residues: Asp-21 to Asp-28, Ser-34 to Asp-40. 751119 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4423 as residues: Asp-1 to Arg-9, Asn-51 to Cys-57, Cys-125 to Leu-137, Cys-153 to Trp-166, Leu-181 to Glu-186, Ser-207 to Thr-212. 756466 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4429 as residues: Ser-1 to Asn-8. 756649 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4429 as residues: Ser-1 to Asn-8. 756649 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4430 as residues: Ala-17 to Leu-23, Gl		ID NO. 4404 as residues: Val-36 to Lys-51, Asn-59 to Asn-76, Val-91
173, Thr-188 to Lys-201, Lys-207 to Ile-226, Leu-234 to Thr-258, Glu-260 to Ile-268, Ser-275 to Lys-286, Val-288 to Glu-299. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4407 as residues: Lys-8 to His-18. 734012 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4411 as residues: Lys-34 to Ser-39. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4412 as residues: His-1 to Gln-6, Glu-19 to Val-26. 739061 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4413 as residues: Asn-7 to Lys-13. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4414 as residues: Pro-10 to Trp-18. 741804 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4416 as residues: Asn-21 to Ser-30, His-37 to Lys-48, Phe-75 to Arg-82. 742220 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4417 as residues: Val-17 to Pro-23, Ser-72 to His-79, Thr-93 to Ile-100, Pro-102 to Asp-108, Asn-111 to Tyr-117, Gly-134 to Lys-141. 744605 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4418 as residues: Asp-1 to Lys-11. 745368 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4420 as residues: Lys-10 to Ser-16, Pro-30 to Arg-37. 750486 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4422 as residues: Asp-21 to Asp-28, Ser-34 to Asp-40. 751119 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4423 as residues: Asp-1 to Arg-9, Asn-51 to Cys-57, Cys-125 to Leu-137, Cys-153 to Trp-166, Leu-181 to Glu-186, Ser-207 to Thr-212. 756466 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4429 as residues: Ser-1 to Asn-8. 756649 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4429 as residues: Ser-1 to Asn-8. 756649 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4430 as residues: Ala-17 to Leu-23, Gl		to Lys-107, Leu-112 to Cys-135, Arg-140 to Lys-150, Pro-157 to Glu-
260 to Ile-268, Ser-275 to Lys-286, Val-288 to Glu-299. 731881 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4407 as residues: Lys-34 to Ser-39. 73503 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4411 as residues: Lys-34 to Ser-39. 735603 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4412 as residues: His-1 to Gln-6, Glu-19 to Val-26. 739061 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4413 as residues: Asn-7 to Lys-13. 741134 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4414 as residues: Pro-10 to Trp-18. 741804 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4416 as residues: Asp-21 to Ser-30, His-37 to Lys-48, Phe-75 to Arg-82. 742220 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4417 as residues: Val-17 to Pro-23, Ser-72 to His-79, Thr-93 to Ile-100, Pro-102 to Asp-108, Asn-111 to Tyr-117, Gly-134 to Lys-141. 744605 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4418 as residues: Asp-1 to Lys-11. 745368 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4420 as residues: Lys-10 to Ser-16, Pro-30 to Arg-37. 750486 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4422 as residues: Asp-21 to Asp-28, Ser-34 to Asp-40. 751119 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4425 as residues: Asp-1 to Arg-9, Asn-51 to Cys-57, Cys-125 to Leu-137, Cys-153 to Trp-166, Leu-181 to Glu-186, Ser-207 to Thr-212. 756466 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4425 as residues: Asp-1 to Arg-9, Asn-51 to Cys-57, Cys-125 to Leu-137, Cys-153 to Trp-166, Leu-181 to Glu-186, Ser-207 to Thr-212. 757213 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4430 as residues: Ala-17 to Leu-23, Gl		173, Thr-188 to Lys-201, Lys-207 to Ile-226, Leu-234 to Thr-258, Glu-
Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4407 as residues: Lys-8 to His-18. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4411 as residues: Lys-34 to Ser-39. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4412 as residues: His-1 to Gln-6, Glu-19 to Val-26. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4413 as residues: Asn-7 to Lys-13. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4414 as residues: Pro-10 to Trp-18. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4416 as residues: Asn-21 to Ser-30, His-37 to Lys-48, Phe-75 to Arg-82. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4416 as residues: Val-17 to Pro-23, Ser-72 to His-79, Thr-93 to Ile-100, Pro-102 to Asp-108, Asn-111 to Tyr-117, Gly-134 to Lys-141. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4418 as residues: Asp-1 to Lys-11. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4420 as residues: Lys-10 to Ser-16, Pro-30 to Arg-37. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4422 as residues: Lys-10 to Ser-16, Pro-30 to Arg-37. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4422 as residues: Sap-1 to Asp-28, Ser-34 to Asp-40. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4423 as residues: Asp-21 to Asp-29, Asn-51 to Cys-57, Cys-125 to Leu-137, Cys-153 to Trp-166, Leu-181 to Glu-186, Ser-207 to Thr-212. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4427 as residues: Ser-1 to Asn-8. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4429 as residues: Ser-1 to Asn-8. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4429 as residues: Ser-1 to Asn-8. Preferred epitopes include those comprising a sequence sho		
ID NO. 4407 as residues: Lys-8 to His-18.	731881	
Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4411 as residues: Lys-34 to Ser-39. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4412 as residues: His-1 to Gin-6, Glu-19 to Val-26. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4413 as residues: Asn-7 to Lys-13. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4414 as residues: Pro-10 to Trp-18. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4416 as residues: Asp-21 to Ser-30, His-37 to Lys-48, Phe-75 to Arg-82. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4417 as residues: Val-17 to Pro-23, Ser-72 to His-79, Thr-93 to IIe-100, Pro-102 to Asp-108, Asn-111 to Tyr-117, Gly-134 to Lys-141. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4418 as residues: Asp-1 to Lys-11. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4420 as residues: Lys-10 to Ser-16, Pro-30 to Arg-37. 750486 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4422 as residues: Asp-21 to Asp-28, Ser-34 to Asp-40. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4422 as residues: Asp-21 to Asp-28, Ser-34 to Asp-40. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4423 as residues: Asp-10 to Gly-13, Gly-18 to Glu-29. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4425 as residues: Ser-1 to Arg-9, Asn-51 to Cys-57, Cys-125 to Leu-137, Cys-153 to Trp-166, Leu-181 to Glu-186, Ser-207 to Thr-212. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4429 as residues: Ser-1 to Asn-8. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4429 as residues: Ser-1 to Asn-8. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4430 as residues: Ser-1 to Asn-8. Preferred epitopes include those comprising a sequ		
ID NO. 4411 as residues: Lys-34 to Ser-39. 735603 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4412 as residues: His-1 to Gln-6, Glu-19 to Val-26. 739061 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4413 as residues: Asn-7 to Lys-13. 741134 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4414 as residues: Pro-10 to Trp-18. 741804 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4416 as residues: Asp-21 to Ser-30, His-37 to Lys-48, Phe-75 to Arg-82. 742220 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4417 as residues: Val-17 to Pro-23, Ser-72 to His-79, Thr-93 to Ile-100, Pro-102 to Asp-108, Asn-111 to Tyr-117, Gly-134 to Lys-141. 744605 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4418 as residues: Asp-1 to Lys-11. 745368 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4420 as residues: Lys-10 to Ser-16, Pro-30 to Arg-37. 750486 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4422 as residues: Asp-21 to Asp-28, Ser-34 to Asp-40. 751119 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4423 as residues: Gly-1 to Gly-13, Gly-18 to Glu-29. 753226 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4425 as residues: Asp-1 to Arg-9, Asn-51 to Cys-57, Cys-125 to Leu-137, Cys-153 to Trp-166, Leu-181 to Glu-186, Ser-207 to Thr-212. 756466 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4429 as residues: Gly-1 to His-10, His-21 to Asp-32. 757213 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4429 as residues: Gly-1 to His-10, His-21 to Asp-32. 757808 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4431 as residues: Ser-23 to Arg-32, Glu-39 to Thr-45, Glu-52 to Lys-57.	734012	
Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4412 as residues: His-1 to Gin-6, Glu-19 to Val-26. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4413 as residues: Asn-7 to Lys-13. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4414 as residues: Pro-10 to Trp-18. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4416 as residues: Asp-21 to Ser-30, His-37 to Lys-48, Phe-75 to Arg-82. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4417 as residues: Val-17 to Pro-23, Ser-72 to His-79, Thr-93 to Ile-100, Pro-102 to Asp-108, Asn-111 to Tyr-117, Gly-134 to Lys-141. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4418 as residues: Asp-1 to Lys-11. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4420 as residues: Lys-10 to Ser-16, Pro-30 to Arg-37. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4422 as residues: Asp-21 to Asp-28, Ser-34 to Asp-40. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4423 as residues: Gly-1 to Gly-13, Gly-18 to Glu-29. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4425 as residues: Asp-1 to Arg-9, Asn-51 to Cys-57, Cys-125 to Leu-137, Cys-153 to Trp-166, Leu-181 to Glu-186, Ser-207 to Thr-212. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4427 as residues: Ser-1 to Asn-8. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4429 as residues: Asp-1 to Arg-9, Asn-51 to Cys-57, Cys-125 to Leu-137, Cys-153 to Trp-166, Leu-181 to Glu-186, Ser-207 to Thr-212. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4429 as residues: Ser-1 to Asn-8. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4430 as residues: Asp-10. His-10, His-21 to Asp-32. Preferred epitopes include those comprising a sequence shown in SEQ ID NO.	754012	
ID NO. 4412 as residues: His-1 to Gin-6, Glu-19 to Val-26.	735603	
739061 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4413 as residues: Asn-7 to Lys-13. 741134 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4414 as residues: Pro-10 to Trp-18. 741804 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4416 as residues: Asp-21 to Ser-30, His-37 to Lys-48, Phe-75 to Arg-82. 742220 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4417 as residues: Val-17 to Pro-23, Ser-72 to His-79, Thr-93 to IIe-100, Pro-102 to Asp-108, Asn-111 to Tyr-117, Gly-134 to Lys-141. 744605 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4418 as residues: Asp-1 to Lys-11. 745368 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4420 as residues: Lys-10 to Ser-16, Pro-30 to Arg-37. 750486 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4422 as residues: Asp-21 to Asp-28, Ser-34 to Asp-40. 751119 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4423 as residues: Gly-1 to Gly-13, Gly-18 to Glu-29. 753226 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4425 as residues: Asp-1 to Arg-9, Asn-51 to Cys-57, Cys-125 to Leu-137, Cys-153 to Trp-166, Leu-181 to Glu-186, Ser-207 to Thr-212. 756466 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4427 as residues: Ser-1 to Asn-8. 756649 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4429 as residues: Gly-1 to His-10, His-21 to Asp-32. 757213 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4430 as residues: Ala-17 to Leu-23, Gly-28 to Gly-42, His-55 to Glu-62, Gly-92 to Ala-100. 757508 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4431 as residues: Ser-23 to Arg-32, Glu-39 to Thr-45, Glu-52 to Lys-57.	/33003	
ID NO. 4413 as residues: Asn-7 to Lys-13.	720061	
741134 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4414 as residues: Pro-10 to Trp-18. 741804 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4416 as residues: Asp-21 to Ser-30, His-37 to Lys-48, Phe-75 to Arg-82. 742220 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4417 as residues: Val-17 to Pro-23, Ser-72 to His-79, Thr-93 to Ile-100, Pro-102 to Asp-108, Asn-111 to Tyr-117, Gly-134 to Lys-141. 744605 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4418 as residues: Asp-1 to Lys-11. 745368 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4420 as residues: Lys-10 to Ser-16, Pro-30 to Arg-37. 750486 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4422 as residues: Asp-21 to Asp-28, Ser-34 to Asp-40. 751119 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4423 as residues: Gly-1 to Gly-13, Gly-18 to Glu-29. 753226 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4423 as residues: Asp-1 to Arg-9, Asn-51 to Cys-57, Cys-125 to Leu-137, Cys-153 to Trp-166, Leu-181 to Glu-186, Ser-207 to Thr-212. 756466 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4429 as residues: Ser-1 to Asn-8. 756649 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4429 as residues: Gly-1 to His-10, His-21 to Asp-32. 757213 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4430 as residues: Ala-17 to Leu-23, Gly-28 to Gly-42, His-55 to Glu-62, Gly-92 to Ala-100. 757508 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4431 as residues: Ser-23 to Arg-32, Glu-39 to Thr-45, Glu-52 to Lys-57. 757980 Preferred epitopes include those comprising a sequence shown in SEQ	/39061	
ID NO. 4414 as residues: Pro-10 to Trp-18.		
Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4416 as residues: Asp-21 to Ser-30, His-37 to Lys-48, Phe-75 to Arg-82. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4417 as residues: Val-17 to Pro-23, Ser-72 to His-79, Thr-93 to Ile-100, Pro-102 to Asp-108, Asn-111 to Tyr-117, Gly-134 to Lys-141. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4418 as residues: Asp-1 to Lys-11. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4420 as residues: Lys-10 to Ser-16, Pro-30 to Arg-37. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4422 as residues: Asp-21 to Asp-28, Ser-34 to Asp-40. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4423 as residues: Gly-1 to Gly-13, Gly-18 to Glu-29. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4425 as residues: Asp-1 to Arg-9, Asn-51 to Cys-57, Cys-125 to Leu-137, Cys-153 to Trp-166, Leu-181 to Glu-186, Ser-207 to Thr-212. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4427 as residues: Ser-1 to Asn-8. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4429 as residues: Ser-1 to Asn-8. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4429 as residues: Ala-17 to Leu-23, Gly-28 to Gly-42, His-55 to Glu-62, Gly-92 to Ala-100. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4430 as residues: Ala-17 to Leu-23, Gly-28 to Gly-42, His-55 to Glu-62, Gly-92 to Ala-100. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4431 as residues: Ser-23 to Arg-32, Glu-39 to Thr-45, Glu-52 to Lys-57. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4431 as residues: Ser-23 to Arg-32, Glu-39 to Thr-45, Glu-52 to Lys-57.	741134	
ID NO. 4416 as residues: Asp-21 to Ser-30, His-37 to Lys-48, Phe-75 to Arg-82. 742220 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4417 as residues: Val-17 to Pro-23, Ser-72 to His-79, Thr-93 to Ile-100, Pro-102 to Asp-108, Asn-111 to Tyr-117, Gly-134 to Lys-141. 744605 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4418 as residues: Asp-1 to Lys-11. 745368 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4420 as residues: Lys-10 to Ser-16, Pro-30 to Arg-37. 750486 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4422 as residues: Asp-21 to Asp-28, Ser-34 to Asp-40. 751119 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4423 as residues: Gly-1 to Gly-13, Gly-18 to Glu-29. 753226 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4425 as residues: Asp-1 to Arg-9, Asn-51 to Cys-57, Cys-125 to Leu-137, Cys-153 to Trp-166, Leu-181 to Glu-186, Ser-207 to Thr-212. 756466 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4427 as residues: Ser-1 to Asn-8. 756649 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4429 as residues: Gly-1 to His-10, His-21 to Asp-32. 757213 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4430 as residues: Ala-17 to Leu-23, Gly-28 to Gly-42, His-55 to Glu-62, Gly-92 to Ala-100. 757508 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4431 as residues: Ser-23 to Arg-32, Glu-39 to Thr-45, Glu-52 to Lys-57. 757980 Preferred epitopes include those comprising a sequence shown in SEQ		
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757508 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4431 as residues: Ser-23 to Arg-32, Glu-39 to Thr-45, Glu-52 to Lys-57. 757980 Preferred epitopes include those comprising a sequence shown in SEQ		· · ·
ID NO. 4431 as residues: Ser-23 to Arg-32, Glu-39 to Thr-45, Glu-52 to Lys-57. 757980 Preferred epitopes include those comprising a sequence shown in SEQ		
Lys-57. 757980 Preferred epitopes include those comprising a sequence shown in SEQ	757508	
757980 Preferred epitopes include those comprising a sequence shown in SEQ		I
ID NO. 4433 as residues: Phe-9 to His-21.	757980	
		ID NO. 4433 as residues: Phe-9 to His-21.

760141	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4434 as residues: Ser-15 to Gly-21, Asp-35 to His-41, Glu-45 to
	Lys-68, Thr-91 to Trp-103, Glu-105 to Gln-116, Asp-124 to Gly-130,
İ	Asp-137 to Thr-147, Glu-162 to Gly-188, Lys-205 to Gly-212, Asn-223
	to Trp-229, Arg-241 to Lys-254.
761491	
101151	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4435 as residues: Gly-55 to Glu-63.
764179	Preferred epitopes include those comprising a sequence shown in SEQ
101175	ID NO. 4438 as residues: Asn-1 to Thr-7.
766961	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4439 as residues: Leu-5 to Glu-16.
768034	Preferred epitopes include those comprising a sequence shown in SEQ
1	ID NO. 4441 as residues: Ser-20 to Lys-29.
769965	Preferred enitones include these commissions and the second secon
1 .0,,,,,	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4442 as residues: Asn-1 to Ser-9, Pro-11 to Cys-38, Pro-41 to
	Val-46, Trp-55 to Ser-62, Pro-73 to Phe-78, Leu-97 to Gln-103, Arg-110 to Gly-116.
771486	
//1460	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4443 as residues: Glu-16 to Lys-21.
772044	Drafarred enitenes in the delay of the last of the las
//2044	Preferred epitopes include those comprising a sequence shown in SEQ
772357	ID NO. 4444 as residues: Ala-11 to Ala-23.
1/233/	Preferred epitopes include those comprising a sequence shown in SEQ
772077	ID NO. 4445 as residues: Phe-61 to Glu-66.
772876	Preferred epitopes include those comprising a sequence shown in SEQ
77.4010	ID NO. 4446 as residues: Arg-80 to Thr-91.
774019	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4447 as residues: Ser-1 to Cys-9, Gln-22 to Gln-28, Gly-41 to
<u> </u>	Gly-4/, Leu-5/ to Arg-66.
774516	Preferred epitopes include those comprising a sequence shown in SEQ
	D NO. 4449 as residues: Leu-41 to Gln-48.
775355	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4450 as residues: Ser-40 to Ala-46.
775367	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4451 as residues: Lys-8 to Lys-28.
775791	Preferred epitopes include those comprising a sequence shown in SEQ
	[ID NO. 4452 as residues: Arg-19 to Asp-29, Asp-81 to Lys-86
778583	Preferred epitopes include those comprising a sequence shown in SEO
	ID NO. 4455 as residues: Thr-10 to Trp-16, Gly-41 to Phe-46, Ser-55 to
	Phe-65.
779588	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4457 as residues: Leu-19 to Lys-26.
781085	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4458 as residues: Ala-57 to Ser-64, Lys-69 to Thr-75.
781366	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4460 as residues: Arg-24 to Pro-35, Gly-72 to His-77.
781376	Preferred enitages include these securities of the preferred enitages include these securities are the preferred enitages include these securities are the preferred enitages include these securities are the preferred enitages include the preferred enitages in the prefe
	Preferred epitopes include those comprising a sequence shown in SEQ
782276	ID NO. 4461 as residues: Pro-39 to Cys-44, Pro-54 to Gly-65.
102210	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4463 as residues: Ile-1 to Gln-9, Arg-27 to Pro-34, Val-36 to

	Pro-60, Lys-86 to Asp-95, Lys-102 to Ser-113, Ser-118 to Asn-130, Asp-132 to Lys-143, Asp-151 to Glu-157.
783413	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4465 as residues: Lys-33 to Val-39.
783668	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4466 as residues: Gly-8 to Leu-17, Leu-27 to Ser-36, Pro-41 to
	Ser-51.
785087	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4468 as residues: Lys-26 to Lys-42.
785465	Preferred epitopes include those comprising a sequence shown in SEQ
/65405	ID NO. 4470 as residues: Gly-6 to Arg-21.
500/0/	
788626	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4471 as residues: Leu-1 to Lys-21, Asp-26 to Asp-34, Ala-85 to
	Тут-90.
788838	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4472 as residues: Ala-14 to Ile-19, Glu-48 to Glu-54, Gln-76 to
	Glu-89.
789419	Preferred epitopes include those comprising a sequence shown in SEQ
/03413	
500(01	ID NO. 4474 as residues: Pro-16 to Asn-22.
789631	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4475 as residues: Thr-10 to Gly-18.
789872	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4476 as residues: Ser-1 to Phe-16, His-36 to Gly-45, Pro-49 to
	Pro-71, Pro-77 to Lys-84.
790190	Preferred epitopes include those comprising a sequence shown in SEQ
,,,,,,	ID NO. 4477 as residues: Ser-41 to Thr-49.
790547	Preferred epitopes include those comprising a sequence shown in SEQ
790347	ID NO. 4478 as residues: Leu-1 to Gln-19, Glu-24 to Pro-31, Lys-36 to
	· · · · · · · · · · · · · · · · · · ·
700557	Cys-45.
792557	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4483 as residues: Lys-51 to Arg-58.
792624	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4484 as residues: Ser-15 to Lys-22, Pro-25 to Gly-47, Glu-55 to
	Thr-64.
793437	Preferred epitopes include those comprising a sequence shown in SEQ
,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,	ID NO. 4485 as residues: Pro-1 to Gly-7, Thr-9 to Phe-18, Ala-32 to
	Trp-45, Pro-53 to Leu-60, Thr-66 to Arg-71.
706022	
796023	Preferred epitopes include those comprising a sequence shown in SEQ
ľ	ID NO. 4488 as residues: Ala-69 to Cys-74, Ile-131 to Glu-136, Gly-
	161 to Asn-169, Leu-174 to Trp-185.
796181	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4489 as residues: Ser-26 to Arg-32, Ala-81 to Cys-87, Pro-118
1	to Lys-126.
797079	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4490 as residues: Phe-2 to Cys-8, Ser-30 to His-36.
797477	Preferred epitopes include those comprising a sequence shown in SEQ
191411	
707:06	ID NO. 4491 as residues: Gly-14 to Leu-24.
797486	Preferred epitopes include those comprising a sequence shown in SEQ
L	ID NO. 4492 as residues: Ser-18 to Gln-25, Pro-35 to Thr-44, Pro-94 to

	Trp-99, Gln-108 to Ser-120, Pro-182 to Gly-187, Pro-192 to Gly-198, Trp-284 to Thr-292.
797747	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4493 as residues: Asn-2 to Ala-11, His-35 to Pro-40.
805448	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4496 as residues: Leu-1 to Tyr-7, Gly-15 to Asn-26.
806690	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4497 as residues: Gly-34 to Trp-43, Trp-48 to Lys-54.
810870	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4498 as residues: Val-12 to Ile-21.
811047	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4499 as residues: Phe-8 to Gly-13, Glu-16 to Asn-34, Ser-179 to Cys-185, Thr-206 to Phe-219.
812745	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4500 as residues: Gly-50 to His-62, Lys-169 to Arg-174, Thr-200 to Asp-206, Leu-208 to Gly-214, Pro-244 to Glu-254, Asp-304 to Gln-310, Gln-318 to Trp-323, Thr-410 to His-415.
812871	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4502 as residues: Ser-22 to Arg-29.
813482	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4503 as residues: Cys-53 to His-65, Glu-71 to Gln-91, Asn-123 to Phe-131, Ala-157 to Pro-171, Gln-197 to Ala-238.
815696	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4504 as residues: Arg-80 to Glu-86, Pro-102 to Thr-110, Pro-113 to Phe-122, Asn-124 to Tyr-131, Thr-149 to Cys-156, Thr-184 to Pro-196, Ser-203 to Cys-215, Gly-226 to Asp-231, Pro-285 to Gly-290.
821335	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4505 as residues: Ser-47 to Cys-59.
827315	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4508 as residues: Asp-29 to Phe-36, Phe-39 to Gly-51.
827740	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4511 as residues: Ile-22 to Lys-28.
828180	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4512 as residues: Glu-38 to Arg-52, Ser-56 to Val-62.
828552	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4513 as residues: Ser-1 to Ser-10, Leu-64 to Asp-69, Gly-102 to Arg-107.
828919	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4515 as residues: Thr-49 to Val-54, Leu-83 to Lys-91, Gly-121 to Thr-130, Asp-165 to Glu-172, Thr-180 to Gly-188.
829084	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4516 as residues: Glu-37 to Trp-47.
829148	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4517 as residues: Pro-33 to Lys-40.
829161	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4518 as residues: Met-5 to Glu-18, Asp-24 to Tyr-30.
830123	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4519 as residues: Ala-20 to Arg-25.
830194	Preferred epitopes include those comprising a sequence shown in SEQ

	ID NO. 4521 as residues: Ala-43 to Lys-51, Glu-66 to Leu-74, His-81 to
	Glu-88, Arg-98 to Ser-105, Gly-111 to Gln-116, Leu-166 to Lys-182,
	Leu-261 to Ala-273, Glu-294 to Arg-302, Glu-335 to Asp-347.
830343	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4524 as residues: Ser-19 to Gly-24, Lys-73 to Leu-94, Ala-101
	to Arg-112, Gly-137 to Ala-143, Glu-160 to Arg-168, Ser-173 to Lys-
	183.
830347	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4525 as residues: Asp-33 to Ala-39.
830382	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4526 as residues: Leu-47 to Val-63, Ser-69 to Ser-76.
830465	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4528 as residues: Pro-1 to Thr-8, Ser-54 to Gln-61, Thr-80 to
	Thr-85, Gln-92 to Tyr-98, Gln-154 to Gln-162, Glu-172 to Ile-177, Val-
	181 to Lys-188, Lys-213 to Asn-225, Ser-234 to Pro-239, Ile-294 to Lys-
	307, Gly-350 to Asn-355.
830498	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4529 as residues: Pro-39 to Asn-47.
830540	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4530 as residues: Leu-31 to Lys-37, Arg-48 to Asn-54.
830586	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4533 as residues: Pro-1 to Gln-15, Arg-33 to Leu-40, Arg-72 to
1	Ser-78, Leu-98 to Asp-103, Phe-116 to Gly-124, Pro-152 to Arg-158,
	Thr-193 to Pro-200, Leu-213 to Phe-219, Asp-229 to Lys-237, Lys-246
	to Lys-258, Arg-275 to Thr-280, Thr-306 to Lys-312, Leu-320 to Arg-
	328, Ala-335 to Asn-340, Gly-342 to Trp-349, Cys-364 to Pro-372.
830693	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4535 as residues: Met-2 to Thr-12, Gln-52 to Glu-67, Glu-72 to
	Val-79, Asn-158 to Arg-165, Met-173 to Gln-180, Glu-200 to Arg-206,
	Ala-220 to Ala-228, Arg-232 to Leu-242, Asp-246 to Gln-254, Thr-260
920722	to Lys-267, Leu-343 to Glu-349. Preferred epitopes include those comprising a sequence shown in SEQ
830723	ID NO. 4537 as residues: Ile-68 to Thr-75, Asp-106 to Asp-117.
830743	Preferred epitopes include those comprising a sequence shown in SEQ
830743	ID NO. 4538 as residues: Pro-11 to Phe-16, Thr-48 to Ser-60.
920904	Preferred epitopes include those comprising a sequence shown in SEQ
830804	ID NO. 4539 as residues: Thr-62 to Gly-70.
920916	Preferred epitopes include those comprising a sequence shown in SEQ
830816	ID NO. 4540 as residues: Thr-51 to Asp-61, Pro-92 to Asn-100, Thr-
	131 to Asn-138, Lys-140 to His-151, Glu-168 to Arg-184, Glu-192 to
	Glu-197, Ala-202 to Leu-212, Tyr-218 to Lys-223, Ala-239 to Leu-246,
	Leu-250 to Gly-256, Pro-289 to Glu-295, Lys-314 to Lys-326, Gln-335
	to Glu-340, Asp-354 to Ser-359.
830829	Preferred epitopes include those comprising a sequence shown in SEQ
030029	ID NO. 4541 as residues: Pro-16 to His-21, Cys-28 to His-35, Val-43 to
	Arg-49, Pro-116 to Tyr-123.
830859	Preferred epitopes include those comprising a sequence shown in SEQ
630639	ID NO. 4542 as residues: Gln-13 to His-28, Pro-73 to Gly-80, Pro-87 to
	Asn-92.
830879	Preferred epitopes include those comprising a sequence shown in SEQ
030079	Treferred epitopes metade mose comprising a sequence shown in 520

	ID NO. 4543 as residues: Cys-34 to Leu-44, Ser-60 to Gly-69, Asp-118
830901	to Gly-123, Cys-148 to Gln-154. Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4544 as residues: Arg-8 to Ser-16, Val-32 to Thr-38, Glu-139 to Lys-145, Arg-224 to Arg-232.
831019	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4545 as residues: Phe-16 to Ser-21.
831057	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4546 as residues: Arg-1 to Gly-14, Thr-19 to Gly-25, Ala-31 to Ala-41, Glu-53 to Ile-62, Val-66 to Glu-75, Ser-103 to Asp-113, Ala-135 to Asp-140.
831099	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4547 as residues: Leu-12 to Gly-18, Leu-93 to Ile-98, Lys-165 to Ser-183, Thr-198 to Lys-211, Glu-232 to Gly-237, Pro-239 to Gly-249, Arg-257 to Asp-278, Cys-292 to Glu-297, Arg-306 to Ser-316, Asp-323 to Asn-331, Glu-347 to Gly-354, Thr-365 to Asn-370, Pro-390 to Thr-396, Asn-420 to Ser-433, Val-440 to Gln-451, His-457 to Asp-465, Phe-533 to Met-538, Ala-540 to Tyr-550, Pro-560 to Lys-565.
831117	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4548 as residues: Lys-50 to Tyr-55.
831163	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4549 as residues: Ser-31 to Arg-40.
831212	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4551 as residues: Arg-34 to Gly-45, Pro-50 to Ala-58.
831234	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4552 as residues: Gly-28 to Pro-33, Gln-66 to Gln-72.
831268	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4554 as residues: Ser-16 to Lys-21.
831307	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4555 as residues: Pro-19 to Ile-26, Ala-43 to Thr-49, Ser-52 to Lys-69, Phe-126 to Arg-134, Pro-153 to Phe-161, Ser-192 to Leu-198, Arg-222 to Thr-229.
831390	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4558 as residues: Trp-50 to Gly-55, Leu-109 to Val-119, Phe-146 to Asp-158, Ser-165 to Trp-172, Phe-192 to Ile-197, Leu-241 to Asp-252, Lys-268 to Pro-273, Ser-310 to Lys-315, Asp-334 to Ala-342, Pro-348 to Tyr-353.
831426	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4559 as residues: Gly-8 to Phe-18, His-26 to Phe-41, Glu-56 to Gly-62, Phe-114 to Lys-126, Asn-198 to Ser-203, Asn-234 to Ile-242, Glu-264 to Pro-270.
831453	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4560 as residues: Tyr-34 to His-42, Leu-44 to Leu-49.
831465	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4561 as residues: Thr-2 to Ser-9, Pro-23 to Ser-28, Phe-55 to Ala-60, Phe-72 to Ile-77, Leu-124 to Gly-136, Glu-138 to Val-144.
831586	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4563 as residues: Gln-14 to Glu-28.
831664	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4564 as residues: Lys-1 to Asp-42, Arg-71 to Ala-76, Gln-138

	to Phe-145, Lys-170 to Thr-178, Cys-186 to Asp-192.
831687	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4565 as residues: Ala-56 to Tyr-63.
831753	Preferred epitopes include those comprising a sequence shown in SEQ
001100	ID NO. 4567 as residues: His-10 to Gly-16, Gly-30 to Phe-36, Ala-41 to
	Lys-47, Phe-63 to Trp-72.
831757	Preferred epitopes include those comprising a sequence shown in SEQ
651757	ID NO. 4568 as residues: Val-81 to Lys-86.
831795	Preferred epitopes include those comprising a sequence shown in SEQ
031/93	ID NO. 4569 as residues: Asn-23 to Pro-28, Arg-36 to Ser-42.
921706	· · · · · · · · · · · · · · · · · · ·
831796	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4570 as residues: Pro-1 to Ser-8.
831880	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4571 as residues: Asp-18 to Ser-24, His-34 to Gly-47.
831899	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4572 as residues: Asp-11 to Trp-16, Pro-37 to Thr-44, Pro-74 to
	Pro-82, Arg-112 to Gln-119, Cys-126 to Arg-138, Arg-199 to Thr-204.
831910	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4573 as residues: Gly-15 to Trp-21, Ser-84 to Leu-93.
831931	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4574 as residues: Asn-29 to Ser-34.
831942	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4575 as residues: Arg-14 to Trp-19, Pro-29 to Gly-37, Cys-51
	to Ala-62, Glu-84 to Glu-91, Ile-101 to Pro-107, Glu-118 to Thr-123,
	Lys-170 to Gln-175, Thr-197 to Lys-228.
832009	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4577 as residues: Leu-17 to Arg-32.
832010	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4578 as residues: Leu-1 to Lys-21, Glu-39 to Cys-47, Lys-49 to
	Gln-61, His-64 to Gly-76, Thr-83 to Lys-90, His-92 to Ile-99.
832093	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4580 as residues: Pro-29 to Tyr-35, Phe-37 to His-42.
832187	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4583 as residues: Glu-11 to Pro-24, Gly-90 to Leu-96, Ser-109
	to Gly-120.
832575	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4588 as residues: Thr-24 to Arg-29, Ala-55 to Tyr-60, Tyr-77 to
	Asp-89, Leu-108 to Gly-115, Thr-142 to Gly-149.
832593	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4589 as residues: Glu-13 to Glu-18.
832597	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4590 as residues: Val-3 to Asp-13.
834890	Preferred epitopes include those comprising a sequence shown in SEQ
33.070	ID NO. 4591 as residues: Arg-8 to Lys-13, Gly-35 to Lys-42, Ala-48 to
	Lys-54, Ala-105 to Leu-110, Gly-150 to Val-157, Phe-164 to Asn-173.
835079	Preferred epitopes include those comprising a sequence shown in SEQ
033013	ID NO. 4592 as residues: Ser-53 to Pro-60.
835456	Preferred epitopes include those comprising a sequence shown in SEQ
655450	ID NO. 4593 as residues: Thr-2 to Asn-10, Ser-72 to Lys-78, Gly-95 to
	1D 140. +333 as residues. Till-2 to Asil-10, Sel-72 to Lys-76, Oly-33 to

	Thr-101, Phe-134 to Ile-147, Lys-163 to Lys-172, Gln-199 to Glu-206,
ĺ	Ala-212 to Trp-224, Lys-230 to His-236, Arg-238 to Glu-244, Asp-249
	to Gly-254, Met-260 to Tyr-266, Arg-272 to Arg-279.
835655	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4594 as residues: Lys-24 to Asn-36, Glu-55 to Asn-60.
836203	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4595 as residues: Pro-43 to Cys-49, Ser-67 to Glu-76, Lys-105
	to Cys-110.
836762	Preferred epitopes include those comprising a sequence shown in SEQ
030702	ID NO. 4507 as residues: Are 262 to Physical Sequence snown in SEQ
838459	ID NO. 4597 as residues: Arg-252 to Phe-260, Ser-315 to Thr-321.
050459	Preferred epitopes include those comprising a sequence shown in SEQ
920262	ID NO. 4600 as residues: Asp-1 to Lys-14.
839262	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4601 as residues: Lys-29 to Asp-36, Gln-98 to Asp-103, Thr-
	120 to Lys-142, Thr-158 to Ser-170, Ile-188 to Glu-194, Leu-217 to
	Gly-223, Tyr-245 to His-252.
839750	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4603 as residues: Gln-27 to Pro-33.
840028	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4604 as residues: Ala-16 to Asn-25, His-32 to Asn-37, Pro-97
	to Ser-103, Pro-114 to Ser-120.
840675	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4606 as residues: Pro-134 to Thr-145.
840708	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4607 as residues: Ala-27 to Ser-36.
840848	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4609 as residues: Arg-77 to Asn-82, Glu-119 to Arg-124, Gln-
	156 to Thr-162, Lys-209 to Lys-215.
840860	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4610 as recidues: The 27 to Apr. 41 Chy. 42 to A1. 50 Chy. 42
	ID NO. 4610 as residues: Ile-27 to Asp-41, Glu-43 to Ala-58, Glu-149
	to Glu-154, Lys-158 to Ile-165, Glu-167 to Gly-189, Glu-242 to Phe-
	247, Arg-259 to Phe-268, Ile-283 to Val-291, Thr-295 to Thr-307, Glu-
841015	328 to Asp-338, Asp-372 to Gly-387.
041013	Preferred epitopes include those comprising a sequence shown in SEQ
941017	ID NO. 4611 as residues: Tyr-17 to Thr-29, Lys-35 to Glu-40.
841017	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4612 as residues: Gln-1 to Trp-19.
841030	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4613 as residues: Ser-23 to Gln-30.
841241	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4614 as residues: Asp-41 to Ile-52, Thr-59 to Lys-64, Glu-75 to
	Asn-89, Thr-99 to Thr-105.
841957	Preferred epitopes include those comprising a sequence shown in SEQ
i	ID NO. 4615 as residues: Gly-7 to Thr-20, Pro-44 to Thr-49, Gln-55 to
}	Gly-61.
846025	
0.3025	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4616 as residues: Gly-8 to Gly-28, Glu-113 to Asn-122, Arg-
	144 to Gly-214, Ala-218 to Gly-232, Arg-243 to Glu-248, Glu-356 to Ser-366.
<u> </u>	SCI-300.

846362	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4617 as residues: His-8 to Gly-18, Phe-66 to Asp-72, Pro-95 to Gly-109, Thr-118 to Ala-126, Gly-128 to Gly-135, Pro-187 to Ser-192, Gly-252 to Arg-258, Asp-270 to Cys-277, Ser-339 to Leu-345, Gly-450 to Ala-468.
846384	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4618 as residues: Gly-3 to Leu-9, Arg-35 to Gly-42, Asp-50 to Thr-55, Ser-98 to Asn-103, Pro-172 to Gly-178, Ser-233 to Pro-243, Ala-289 to Gly-294, Thr-302 to Tyr-309, Glu-341 to Trp-347, Pro-349 to Val-359, Pro-414 to Thr-422, Arg-438 to Glu-443, Gln-507 to Thr-518.
846750	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4619 as residues: Thr-27 to Arg-32, Gly-63 to Gly-71, Ile-95 to Gly-101, Asn-108 to Ser-115.
847598	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4621 as residues: Ser-1 to Thr-27.
848119	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4622 as residues: Pro-5 to Lys-10, Ser-29 to Lys-42, Arg-54 to Arg-66.
848746	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4623 as residues: Pro-61 to Asp-68, Arg-88 to Asp-93.
849084	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4624 as residues: Gly-1 to Pro-8, Ala-48 to Tyr-53, Lys-55 to Arg-62, Glu-67 to Leu-75.
849114	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4625 as residues: Asn-30 to Leu-36, Trp-51 to Phe-56, Pro-62 to Trp-68, Gln-98 to Ser-114, Ile-128 to His-134, Pro-146 to His-151, Asp-153 to Tyr-171, Asp-193 to Trp-198, Pro-222 to Thr-234, Ile-237 to Thr-260, Ile-285 to Gly-296, Arg-301 to Gln-308, Val-311 to Asp-328.
849155	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4627 as residues: Pro-6 to Lys-21, Ala-26 to Val-34, Lys-37 to Ser-46, Phe-73 to Val-81, Pro-86 to Arg-92, Gly-101 to Ser-108, Thr-172 to Pro-178, Met-244 to Lys-255.
849159	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4628 as residues: Thr-28 to Ala-33, Asn-93 to Trp-103, Ile-122 to Pro-130, His-132 to Ile-138.
849244	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4629 as residues: Gln-189 to Glu-196, Glu-206 to Pro-211, Ser-226 to Ile-233, Lys-244 to Ser-253.
849254	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4630 as residues: Ala-5 to Cys-11, Cys-14 to Gly-25, Tyr-32 to Gln-38, Glu-62 to Leu-78, Asp-91 to Tyr-102.
849301	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4631 as residues: Ser-37 to Asp-43, Lys-266 to Ser-272, Glu-304 to Thr-318, Leu-345 to Ser-359, Gln-423 to Ala-439.
849317	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4632 as residues: Pro-42 to Trp-47, Arg-49 to Glu-55, Val-62 to Glu-67, Leu-75 to Leu-90, Leu-102 to Gln-107, Ile-154 to Asp-161.
849332	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4633 as residues: Gln-31 to Ser-38, Gly-60 to Arg-65, Thr-148

	to Thr-155, Cys-180 to Cys-189, Val-224 to Pro-232, Leu-250 to Gln- 255.
849422	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4634 as residues: Arg-9 to Arg-14.
849492	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4636 as residues: Ser-5 to Arg-11.
849534	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4637 as residues: Met-8 to His-14.
849565	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4638 as residues: Gly-59 to Ala-67.
849583	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4639 as residues: Pro-13 to Pro-18, Pro-24 to Leu-32, Glu-51 to His-59, Leu-83 to Trp-91, Thr-113 to Gln-120, Pro-133 to Asp-138, Arg-141 to Gln-146, Arg-151 to Ser-156, Tyr-160 to Cys-175, Asn-183 to Asn-188, Trp-221 to Lys-231, Ser-271 to Arg-283, Phe-345 to Gly-350, Ser-381 to Asp-386, Gly-417 to Ser-422, Tyr-462 to Asn-471, Glu-505 to Leu-533, Ser-555 to Asp-561, Thr-566 to His-576, Ser-582 to Gln-587.
849589	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4640 as residues: Ser-16 to Val-25, His-105 to Lys-125, Tyr-147 to Ser-155.
849658	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4641 as residues: Ser-1 to Ser-7.
849666	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4642 as residues: Glu-12 to Met-22.
849679	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4643 as residues: Lys-208 to Asp-214, Glu-278 to Gln-289, Glu-296 to Arg-303, Lys-358 to Leu-364.
849741	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4644 as residues: Arg-30 to His-40.
849783	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4645 as residues: Arg-1 to Pro-14, Gln-47 to Cys-52, Asn-57 to Pro-63, Ser-277 to Lys-282, Leu-326 to Ser-332.
850211	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4646 as residues: Asn-8 to Asn-13.
850254	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4647 as residues: Asn-1 to Arg-6.
850264	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4648 as residues: Ala-33 to Gly-47, Glu-73 to Lys-78, Ser-111 to Asp-126, Gln-139 to Ala-147, Cys-206 to Gly-211, Ser-218 to Asn-225, Leu-237 to Pro-242, Arg-277 to Leu-282, Lys-284 to Lys-291, Ala-357 to Asn-363, Asn-380 to Leu-387, His-475 to Arg-489, Pro-494 to Lys-515.
850273	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4649 as residues: Pro-31 to Lys-38.
850371	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4650 as residues: Lys-32 to Thr-38.
850859	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4651 as residues: Phe-18 to Lys-24, Pro-53 to Lys-75, Tyr-115 to Asp-124, Lys-130 to Leu-137.

851066	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4652 as residues: Pro-6 to Asp-12, Arg-28 to Thr-37, Ile-50 to
	Lys-59, Ala-63 to Gly-70, Pro-89 to Tyr-96, Ser-103 to Ile-111, Thr-114
	to Phe-121, Asp-141 to Pro-147, Arg-162 to Thr-172.
851217	Preferred epitopes include those comprising a sequence shown in SEQ
<u> </u>	ID NO. 4653 as residues: Gln-24 to Asp-36, Ser-54 to Thr-65.
852170	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4654 as residues: Leu-13 to Glu-26.
852387	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4655 as residues: Ala-37 to Thr-43.
852812	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4656 as residues: Pro-27 to Pro-33, Asp-92 to Gly-99, Asp-109
	to Lys-115, Pro-117 to Trp-130, Phe-208 to Thr-215, Ile-219 to Lys-231,
	Arg-251 to Asp-257.
853175	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4657 as residues: Gln-21 to Ser-31, Tyr-74 to Gln-81, Leu-115
	to Arg-121.
854063	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4659 as residues: Pro-3 to Gly-43.
854073	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4660 as residues: Glu-13 to Val-19, Gln-32 to Met-40, Asp-49
	to Arg-54, Leu-74 to Ser-86.
854987	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4661 as residues: Arg-1 to Arg-12.
855130	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4662 as residues: Glu-64 to Tyr-69.
856227	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4663 as residues: Pro-18 to Arg-35, Ala-42 to Gly-54, His-69 to
	Gln-76, Asp-105 to Arg-110, Arg-121 to Asp-126, Pro-150 to Gln-160.
856243	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4664 as residues: Ala-1 to Ala-8, Lys-78 to Met-86, Arg-126 to
055054	Lys-137.
856354	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4665 as residues: Thr-21 to Thr-33.
858178	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4670 as residues: Gly-2 to Gln-8, Lys-68 to Gln-76, Pro-200 to
	Gly-208, Ser-246 to Gly-257, Gly-280 to Gly-289, Ala-302 to Gly-308,
	Gly-319 to Asn-331, Leu-352 to Ser-361, Glu-378 to Glu-399, Ala-401
252526	to His-414.
858606	Preferred epitopes include those comprising a sequence shown in SEQ
050004	ID NO. 4671 as residues: Trp-86 to Pro-91.
858894	Preferred epitopes include those comprising a sequence shown in SEQ
050050	ID NO. 4672 as residues: Lys-1 to Ser-9.
858958	Preferred epitopes include those comprising a sequence shown in SEQ
950171	ID NO. 4674 as residues: Pro-19 to Ala-25.
859171	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4675 as residues: Lys-12 to Val-18, Leu-32 to Ser-47, Glu-55 to
	Asp-66, Glu-94 to Glu-109, Val-115 to Ile-127, Asp-166 to Ser-177, Lys-213 to Glu-225, Glu-241 to Lys-264, Met-322 to Phe-343, Asn-371
	to Glu-379, Ala-396 to Ser-407, Ser-415 to Pro-422, Pro-435 to Pro-440,

	Ile-459 to Gln-466, Phe-471 to Phe-476.
859352	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4676 as residues: Thr-11 to Thr-21.
859354	Preferred epitopes include those comprising a sequence shown in SEQ
j	ID NO. 4677 as residues: Arg-60 to Pro-70, Ser-138 to Ser-145, Cys-
	157 to Lys-163, Pro-204 to Thr-211, Val-213 to Ser-219, Thr-224 to
	Thr-230, Pro-297 to Asp-302, Ile-332 to Glu-339, Glu-385 to Ser-390.
859702	Preferred epitopes include those comprising a sequence shown in SEQ
26222	ID NO. 4678 as residues: Lys-7 to Arg-26.
860915	Preferred epitopes include those comprising a sequence shown in SEQ
861200	ID NO. 4679 as residues: Gln-50 to Gly-56.
861209	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4680 as residues: Leu-6 to Thr-15, Pro-85 to Asp-90, Thr-98 to
961524	Pro-104.
861534	Preferred epitopes include those comprising a sequence shown in SEQ
061607	ID NO. 4681 as residues: Arg-24 to Ser-30.
861697	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4682 as residues: Gly-8 to Trp-16, Asn-22 to Phe-28, Phe-68 to
	Arg-75, Ser-93 to Ser-101, Glu-114 to Ile-126, Pro-134 to Phe-143, Gly-
	165 to Gly-176, Lys-191 to Glu-201, Thr-218 to Lys-227, Tyr-289 to Gln-296.
861826	
301020	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4683 as residues: Gly-17 to Pro-23.
861909	Preferred epitopes include those comprising a sequence shown in SEQ
001707	ID NO. 4684 as residues: His-13 to Cys-20, Glu-83 to Cys-93, Pro-131
	to Asp-137, Cys-142 to Asn-148, Pro-150 to Gln-155, Pro-160 to Gly-
	166, Ser-194 to Gly-206, Thr-251 to Ser-258, Gly-267 to Asp-272, Lys-
	286 to Gly-299, Gln-353 to Leu-366, Thr-368 to Gln-381, Gln-387 to
	His-397, Glu-404 to Ala-410, Phe-412 to Ala-418, Phe-424 to Ala-439.
862237	Preferred epitopes include those comprising a sequence shown in SEQ
L	ID NO. 4687 as residues: Cys-20 to Val-27.
862285	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4689 as residues: Ala-26 to Gln-32.
862456	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4691 as residues: Pro-20 to Gly-26, Glu-66 to Trp-76.
862486	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4692 as residues: Cys-36 to Pro-44, His-145 to Asn-151, Asp-
	186 to Glu-195, Glu-271 to Ile-281, Asp-296 to Pro-302.
863865	Preferred epitopes include those comprising a sequence shown in SEO
	ID NO. 4694 as residues: Gly-1 to Pro-6, Leu-17 to Ala-22, Phe-40 to
	Ala-45.
863944	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4695 as residues: Glu-102 to Asp-111, Glu-144 to Val-149,
	Tyr-169 to Lys-180, Arg-239 to Arg-245, Gln-247 to Asp-253, Gly-266
-	to Asn-278.
864428	Preferred epitopes include those comprising a sequence shown in SEQ
1	ID NO. 4696 as residues: Thr-1 to Leu-11, Arg-26 to Gly-41, Arg-81 to
	Asp-91, Asp-144 to Thr-159, Asn-170 to Ala-178, Glu-180 to Lys-191,
065011	Cys-249 to Trp-255.
865044	Preferred epitopes include those comprising a sequence shown in SEQ

1800

	ID NO. 4699 as residues: Thr-17 to Gly-34, Pro-66 to Gly-71, Pro-73 to Val-78.
865421	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4701 as residues: Ala-10 to Glu-16.
866287	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4702 as residues: Val-1 to Leu-6.
866300	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4703 as residues: Thr-28 to Trp-35.
867388	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4707 as residues: Ser-39 to Phe-56, Asp-77 to Arg-84, Glu-103 to Lys-129, Lys-134 to Lys-143, Pro-219 to Gly-227, His-289 to Glu-297, Ala-353 to Arg-360, Pro-409 to Tyr-423, His-433 to Thr-441, Phe-445 to Pro-453, Gln-480 to Leu-488, Pro-526 to Thr-540.
867842	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4708 as residues: Leu-38 to His-44, Leu-46 to Gln-55, Leu-65 to Gln-70, Ile-80 to Arg-88.
867923	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4709 as residues: Leu-17 to Leu-23, Gln-51 to Thr-57.
868035	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4710 as residues: Ser-8 to Pro-13, Pro-21 to Ser-33.
868135	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4711 as residues: Glu-27 to Arg-32, Glu-86 to Gly-93, Ala-117 to Glu-127, Glu-148 to Asn-154, Asp-163 to Ser-174.
868173	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4712 as residues: Thr-6 to Asn-14, Pro-19 to Lys-41.
868224	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4713 as residues: Glu-21 to Glu-31, Arg-37 to Ser-45, Asn-47 to Gly-53, Pro-64 to Arg-70, Ser-97 to Tyr-102, Asp-110 to Val-116.
868655	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4714 as residues: Phe-5 to Ser-21, Ser-24 to Ser-32, Ser-40 to Ser-64, Leu-73 to Glu-81, Pro-122 to Leu-130, Glu-186 to Leu-193, Leu-204 to Trp-213, Ser-278 to Ala-285, Glu-376 to Asp-384, Phe-401 to Val-407.
869698	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4715 as residues: Asp-1 to Ser-6, Glu-16 to Ser-26, Lys-66 to Pro-76, Leu-93 to Arg-99, Val-153 to Lys-164, Glu-177 to Asp-183, Ser-188 to Leu-193, Arg-210 to Ser-220, Thr-229 to Ser-244, Pro-283 to Phe-297.
870190	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4716 as residues: Arg-112 to Lys-118, Gln-168 to His-175.
870349	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4717 as residues: Thr-34 to Ala-39, Ser-42 to Arg-47.
870522	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4719 as residues: Asn-32 to Gly-39, Gly-116 to Lys-124.
870896	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4720 as residues: Leu-21 to Gly-30, Arg-41 to Cys-49, Arg-57 to Phe-62.
871071	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4721 as residues: Arg-1 to Cys-13, Lys-26 to Ile-34.

871225	Preferred epitopes include those comprising a sequence shown in SEQ
071 400	10 NO. 4722 as residues: Pro-23 to Gly-36. Arg-77 to Ile-84
871428	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4723 as residues: Gly-6 to Pro-11.
871498	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4724 as residues: Arg-12 to Ser-18.
871732	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4725 as residues: Ser-56 to Thr-62.
871756	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4726 as residues: Ser-31 to Gly-38.
871821	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4727 as residues: Tyr-25 to Lys-30, Lys-36 to Ile-43, Lys-52 to Gln-69, Glu-76 to Asp-81, Arg-92 to Trp-104, Leu-120 to Lys-126, Ser-129 to Ser-135, Ser-139 to Thr-156, Pro-165 to Glu-178, Ser-181 to Thr-186, Tyr-196 to Lys-201, Cys-225 to Lys-230, Glu-234 to Glu-242.
872354	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4729 as residues: Thr-33 to Lys-43, Lys-81 to Ser-100.
872535	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4730 as residues: Ser-33 to Gly-41, Asn-66 to Asp-73, Cys-136 to Gly-141, Met-187 to Thr-193.
872551	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4731 as residues: Cys-1 to Cys-7, Asp-12 to Arg-27, Pro-49 to Tyr-59, Leu-157 to Leu-163, Ser-243 to Thr-248, Thr-349 to Ser-362, Phe-376 to Ser-385.
872640	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4732 as residues: Tyr-1 to Asp-8, Tyr-33 to Gly-39, Glu-57 to Glu-64, Ser-74 to Val-82, Lys-203 to Arg-214, Gln-229 to Pro-235, Gln-310 to Ala-317, Glu-326 to Asn-331, Gly-366 to Asn-372, Leu-392 to Asn-403, Ala-459 to Gln-466, Asp-494 to His-502, Pro-514 to Leu-522, Glu-614 to Leu-621, Asn-642 to His-651.
872802	Preferred epitopes include those comprising a sequence shown in SEO
872852	ID NO. 4734 as residues: Ser-1 to Gly-8, Arg-30 to Trp-37. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4735 as residues: Arg-1 to Gln-7, Arg-22 to Arg-28, Gln-93 to Glu-100.
874307	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4739 as residues: Tyr-1 to Glu-6.
874309	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4741 as residues: Ser-2 to Val-13, Lys-59 to Ser-77.
874310	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4742 as residues: Thr-25 to Thr-31.
874320	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4743 as residues: Ser-1 to Ala-7, Ala-26 to Gly-35, Gly-53 to Phe-59, Arg-67 to Arg-84.
874325	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4744 as residues: Arg-1 to Leu-7, Ser-13 to Val-20, Leu-38 to Glu-44, Leu-79 to Gly-84, Thr-92 to Ala-100, Pro-110 to Ser-119.
874327	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4745 as residues: Asp-45 to Thr-51, Leu-55 to Gly-63, Asp-88 to Phe-97, Gly-185 to Trp-200, Gly-214 to Ser-222, Thr-239 to Val-246.

874329	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4747 as residues: Glu-10 to Ala-16, Asp-32 to His-37.
874348	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4750 as residues: Asn-10 to Thr-15.
874349	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4751 as residues: Pro-1 to Ala-7, Asp-38 to Val-54.
874350	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4752 as residues: Ser-35 to Glu-46, Lys-89 to Asp-94.
874358	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4754 as residues: Phe-34 to Lys-45, Asn-122 to Ser-127, Asp-
	160 to Lys-165.
874362	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4755 as residues: Ile-1 to Ser-12, His-35 to Glu-47, Glu-55 to
	Ser-71, Gly-74 to Ser-82, Ala-97 to Ser-139, Lys-153 to Arg-166, Arg-
ļ	171 to Leu-180, Asp-304 to Gly-309, Glu-373 to Glu-378, Ser-495 to
	Tyr-500.
874368	Preferred epitopes include those comprising a sequence shown in SEQ
0/4300	
	ID NO. 4756 as residues: Ala-14 to Pro-20, Thr-26 to Asn-32, Lys-55
	to Ala-61.
874370	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4758 as residues: Arg-48 to Tyr-55, Tyr-64 to Gly-76.
874372	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4759 as residues: Ala-1 to Gly-16, Lys-33 to Thr-44, Leu-52 to
	Asp-57, Gln-69 to Phe-78, Gly-91 to Cys-104.
874396	Preferred epitopes include those comprising a sequence shown in SEQ
074570	ID NO. 4760 as residues: Leu-39 to Ser-44.
874399	Preferred epitopes include those comprising a sequence shown in SEQ
0/4399	ID NO. 4761 as residues: Pro-36 to Glu-46, Asn-151 to Asn-170, Tyr-
	175 to Thr-180, Glu-182 to Glu-190, Thr-202 to Glu-212, Arg-238 to
	Ser-245, Pro-292 to Gly-302.
874401	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4763 as residues: Gly-10 to Gly-19, Lys-44 to Arg-61, Leu-112
	to Lys-117.
874403	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4764 as residues: Phe-20 to Lys-27, Lys-66 to Arg-82.
874413	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4768 as residues: Phe-1 to Asp-11.
874414	Preferred epitopes include those comprising a sequence shown in SEQ
0,4414	ID NO. 4769 as residues: Ser-54 to Gly-59, Asp-63 to Lys-71.
974416	
874416	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4770 as residues: Thr-7 to Ser-14, Pro-28 to Asp-36.
874417	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4771 as residues: Tyr-16 to Ala-26, Ser-43 to Asp-54.
874423	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4773 as residues: Lys-1 to Gly-8, Ser-55 to Leu-60.
874427	Preferred epitopes include those comprising a sequence shown in SEQ
]	ID NO. 4776 as residues: Tyr-64 to Thr-70.
874435	Preferred epitopes include those comprising a sequence shown in SEQ
014433	
	ID NO. 4780 as residues: Pro-77 to Lys-95.

874437	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4782 as residues: Glu-15 to Glu-29, Ala-43 to Asp-49, Ile-53 to
}	Asp-65, Lys-86 to Pro-94, Val-102 to Gly-121, Asp-160 to Ser-165,
	Asn-234 to Lys-241, Glu-309 to Leu-321, Lys-368 to Ala-377, Thr-382
07110	to Asp-400, Ser-407 to Asn-415, Asp-417 to Leu-448.
874438	Preferred epitopes include those comprising a sequence shown in SEQ
974447	ID NO. 4783 as residues: Pro-19 to Leu-28, Pro-44 to Ser-60.
874447	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4784 as residues: Pro-1 to His-6.
874449	
0/4449	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4785 as residues: Glu-10 to Gly-20, Lys-41 to Met-46, Leu-60
	to Gln-70.
874455	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4787 as residues: Ile-7 to Lys-15.
874459	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4789 as residues: Tyr-1 to Gly-14, Arg-33 to Pro-41, Pro-58 to
	Asp-66.
874468	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4793 as residues: Thr-10 to Arg-15.
874469	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4794 as residues: Gln-19 to Lys-26.
874470	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4795 as residues: Arg-3 to Gly-18, Pro-73 to Glu-86, Ser-104 to
	Pro-117, Gln-143 to Arg-150, Asp-158 to Arg-174, Leu-197 to Ser-222,
974472	Ala-235 to Glu-256, Arg-296 to Arg-309.
874473	Preferred epitopes include those comprising a sequence shown in SEQ
874480	ID NO. 4797 as residues: Ser-28 to Arg-37, Arg-83 to Gln-97.
074400	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4801 as residues: Lys-2 to Gly-8, Pro-54 to Asn-65.
874482	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4803 as residues: Lys-52 to Asn-60.
874484	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4804 as residues: Lys-24 to Ser-38.
874486	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4806 as residues: Trp-1 to Pro-10.
874492	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4807 as residues: Arg-33 to Cys-44.
874495	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4808 as residues: Asp-17 to Val-23, Asp-35 to Trp-40, Phe-63
	to Arg-68, Ala-150 to Thr-156.
874498	Preferred epitopes include those comprising a sequence shown in SEQ
974400	ID NO. 4809 as residues: Ala-37 to Asn-42, Ala-94 to Glu-106.
874499	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4810 as residues: Met-3 to Pro-10, Pro-18 to Arg-23, Pro-62 to
974502	Gly-69.
874503	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4811 as residues: Gln-10 to Glu-21, Ser-28 to Arg-33, Glu-107
874504	to Leu-113, Glu-126 to Ser-133.
0/4304	Preferred epitopes include those comprising a sequence shown in SEQ

	ID NO. 4812 as residues: Pro-53 to Gly-65, Ala-74 to Lys-96, Lys-107 to Lys-116.
874506	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4814 as residues: Ile-81 to Arg-91.
874518	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4816 as residues: Pro-16 to Ser-24, Thr-34 to Pro-39.
874519	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4817 as residues: Asp-19 to Glu-32, Glu-43 to Glu-80.
874522	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4818 as residues: Pro-6 to Pro-12.
874524	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4819 as residues: Asp-16 to Val-21, Leu-33 to Asp-50.
874527	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4820 as residues: Val-1 to Thr-11, Lys-60 to His-73, Met-84 to Gln-99, Thr-119 to Asp-126.
874528	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4821 as residues: Pro-14 to Arg-23, Ala-171 to Ser-178.
874529	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4822 as residues: Pro-7 to Arg-15.
874545	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4830 as residues: Gly-1 to Asp-6.
874550	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4832 as residues: Arg-20 to Lys-28, Leu-40 to Ala-45, Lys-76 to Ser-81, Leu-106 to Lys-111.
874552	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4834 as residues: Ser-70 to Gly-76.
874553	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4835 as residues: Lys-70 to His-78, Lys-149 to Asn-154, Gly-209 to Leu-217, Lys-248 to Val-255, lle-259 to Arg-264, Arg-280 to Ala-287.
874556	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4836 as residues: Pro-73 to Ala-78, Ala-95 to Trp-106, Ala-108 to Gly-121, Lys-132 to Asn-142, Glu-163 to Arg-173, Ser-189 to Glu-194, Val-213 to Leu-229, Gln-244 to Asn-260.
874559	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4837 as residues: Thr-47 to Val-63, Arg-90 to Tyr-102, Val-179 to Pro-187, Asp-189 to Gln-200.
874560	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4838 as residues: Arg-222 to Gly-236, Ser-242 to Ile-250, Leu-254 to Ser-260, Glu-277 to Ser-283.
874561	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4839 as residues: Arg-29 to Gln-45.
874562	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4840 as residues: Pro-65 to Val-75, Pro-101 to Ala-131, Pro-143 to Cys-155, Ser-167 to Pro-179, Thr-205 to Cys-216, Arg-218 to His-236, Gln-241 to Asp-267.
874563	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4841 as residues: Ala-1 to Lys-8.
874564	Preferred epitopes include those comprising a sequence shown in SEQ

	ID NO. 4842 as residues: Pro-1 to Cys-8, Glu-48 to His-58, Ser-72 to Glu-78.
874567	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4844 as residues: Met-46 to Leu-55, Leu-93 to Lys-115, Leu-
1	169 to Gly-187, Glu-213 to Gly-219, Lys-224 to Glu-229, Ser-294 to
	Cys-300, Gln-319 to Leu-328, Ser-345 to Asp-350, Pro-380 to Thr-385,
-	Tyr-387 to Val-393.
874570	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4846 as residues: Pro-3 to Phe-14, Arg-16 to Trp-22, Ser-62 to
	Leu-74, Asp-86 to Ser-92, Gly-102 to Ser-111, Val-113 to Ser-118.
874571	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4847 as residues: Asp-49 to Asp-59, Asp-110 to Ile-115, Trp-
}	137 to Ser-144.
874573	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4848 as residues: Pro-11 to Ala-35, Phe-47 to Glu-54, Glu-78 to
	Gly-83, Gln-94 to Ser-106, Ser-114 to Val-120.
874577	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4849 as residues: Leu-1 to Leu-6, Lys-26 to Asp-44, His-50 to
	Gly-58, Ala-102 to Thr-107.
874580	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4851 as residues: Arg-1 to Val-8, Lys-30 to Tyr-36, Tyr-92 to
	Gly-101, Lys-116 to Lys-125, Asp-140 to Gly-145, Pro-147 to Ser-167,
1	Ser-170 to Ser-191, Ser-193 to Ile-199, Leu-203 to Arg-215, Ser-220 to
	Glu-231.
874581	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4852 as residues: Leu-1 to His-8, Pro-74 to Pro-84.
874590	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4855 as residues: Arg-1 to Asn-13, Pro-34 to Pro-41, Val-77 to
	Thr-84.
874592	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4856 as residues: Val-1 to His-27, Gly-33 to Trp-58, Pro-99 to
	Cys-105.
874594	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4857 as residues: Lys-18 to Gln-27, Leu-41 to Leu-46.
874601	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4859 as residues: Thr-6 to Gly-14, Gly-20 to Ala-26, Pro-31 to
	Met-37, Arg-49 to Ser-64, Pro-70 to His-79.
874605	Preferred epitopes include those comprising a sequence shown in SEQ
ľ	ID NO. 4861 as residues: Val-5 to Gly-11, Ser-43 to Lys-53, Glu-61 to
	Thr-68, Thr-99 to Ala-104, Tyr-106 to Asp-120, Asn-139 to Leu-148,
	Thr-169 to Thr-174, Asn-196 to Asn-202, Asn-223 to Glu-231, Glu-241
	to Tyr-247, Ser-265 to Thr-270, Thr-277 to Cys-286, Leu-292 to Asp-
07//07	298, Asn-347 to Thr-352, Thr-361 to Gly-366, Asn-373 to Thr-383.
874607	Preferred epitopes include those comprising a sequence shown in SEQ
074600	ID NO. 4862 as residues: Pro-1 to Arg-10.
874608	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4863 as residues: Pro-3 to Arg-8, Gly-34 to Thr-53, Asp-60 to
074600	Ser-65, Phe-76 to Lys-81.
874609	Preferred epitopes include those comprising a sequence shown in SEQ
L	ID NO. 4864 as residues: Arg-6 to Arg-13, Phe-25 to Asn-32, Phe-47 to

	Glu-56, Lys-108 to Ala-122.
874610	Preferred epitopes include those comprising a sequence shown in SEQ
074010	ID NO. 4865 as residues: Pro-31 to Trp-39, Pro-101 to Lys-110, Tyr-
	130 to Ala-137, Val-145 to Lys-154, Pro-174 to Gly-179, Phe-194 to
	Asn-202, Glu-224 to Gly-240, Thr-259 to Gln-264, Arg-287 to Ser-293,
	Cys-301 to Gln-307.
874611	Preferred epitopes include those comprising a sequence shown in SEQ
8/4011	ID NO. 4866 as residues: Lys-1 to Gly-6, Asp-13 to Glu-27.
874615	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4869 as residues: Pro-13 to Cys-19.
874618	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4870 as residues: Arg-10 to Cys-15, Phe-30 to Pro-36, Arg-53
	to Ser-59, Thr-66 to Ser-79.
874619	Preferred epitopes include those comprising a sequence shown in SEQ
0,1015	ID NO. 4871 as residues: Ala-1 to Pro-7.
874621	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4873 as residues: Glu-4 to Gly-12, Thr-21 to Gln-27, Pro-40 to
	Ser-47, Pro-50 to Ser-61, Val-101 to Cys-107, Lys-138 to Gly-147, Gln-
	150 to Тут-156, Lys-169 to Thr-174.
874622	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4874 as residues: Gln-31 to Lys-39, His-55 to Asp-60.
874623	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4875 as residues: Arg-7 to His-24, Pro-27 to Gly-33.
874624	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4876 as residues: Gln-12 to Ser-22.
874626	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4878 as residues: Leu-4 to Gly-11, Pro-60 to Gln-65.
874628	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4879 as residues: Pro-13 to Thr-20, His-24 to Gly-34, Glu-36 to
	His-42.
874631	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4881 as residues: Lys-14 to Glu-23, Glu-30 to Ser-43, Ser-45 to
	His-54, Thr-66 to Tyr-71, Pro-75 to Asp-80, Ile-98 to Thr-120, Glu-125
	to Lys-133, Leu-146 to Ala-152, Ala-170 to Ile-176, Asp-180 to Cys-
	200.
874632	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4882 as residues: His-45 to Gly-50.
874635	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4883 as residues: Pro-1 to Pro-7, Leu-19 to Gly-26, Glu-72 to
	Asp-78, Lys-93 to Glu-103, Gln-152 to Gly-159, Gln-181 to Asp-190,
	Phe-232 to Val-237, Asn-282 to Thr-287, Pro-289 to Pro-295, His-341
	to Asp-351, Cys-378 to Glu-383, Gln-448 to Gly-453, Ser-518 to His-
	524, Pro-536 to Glu-541.
874636	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4884 as residues: Glu-1 to Tyr-6, Pro-39 to Asp-46.
874639	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4886 as residues: Pro-7 to Gly-29, Ser-36 to Ala-41, Pro-43 to
	Asp-54, Pro-59 to Leu-64, Gln-70 to Ile-75, Glu-85 to Lys-94.
874642	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4888 as residues: His-8 to Gly-18, Gly-26 to Asp-38.

874644	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4889 as residues: Ser-4 to Leu-10, Thr-25 to Gly-35.
074645	
874645	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4890 as residues: Glu-69 to Thr-75.
874650	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4892 as residues: Glu-2 to Glu-14.
874651	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4893 as residues: Arg-1 to His-9.
874652	Professed enitenes include the annual include the second s
	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4894 as residues: Ser-40 to Asn-45.
874653	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4895 as residues: Thr-1 to Ser-10, Arg-24 to Trp-51, Leu-62 to
	Gly-67, Pro-72 to Gly-81, Pro-98 to Gly-103.
874655	Preferred epitopes include those comprising a sequence shown in SEQ
0,1055	ID NO 4907 as residues. Clu Des Con 14 Ser 29 et 62 47 Th.
	ID NO. 4897 as residues: Glu-9 to Cys-14, Ser-38 to Ser-47, Tyr-52 to
074660	Lys-61, His-68 to Lys-78, Lys-93 to Gly-101.
874660	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4902 as residues: Leu-13 to Glu-18.
874665	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4904 as residues: Arg-9 to Arg-18, Leu-28 to Phe-36, Pro-49 to
	Arg-56, His-85 to Asn-103.
874667	Preferred epitopes include those comprising a sequence shown in SEQ
074007	ID NO. 4905 as residues: Leu-47 to Thr-53, Ala-60 to Ser-66.
874670	D 100. 4903 as residues. Leu-47 to 1111-35, Ala-00 to Ser-00.
8/46/0	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4906 as residues: Lys-1 to Leu-6, Pro-9 to Gly-17, Tyr-19 to
	Glu-25, Arg-30 to Leu-39.
874671	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4907 as residues: Val-5 to Ile-10, Glu-26 to Asp-35, Pro-70 to
	Pro-80, Tyr-90 to Glu-96.
874673	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4909 as residues: Ser-53 to Ser-63.
874675	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4910 as residues: Ser-33 to Ala-48.
874678	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4911 as residues: Lys-1 to Ser-12.
874679	
6/40/9	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4912 as residues: Arg-1 to Glu-7, Leu-21 to Lys-32, His-56 to
	Cys-64.
874680	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4913 as residues: Glu-8 to Arg-14, Ile-49 to His-59, Leu-86 to
	Cys-94.
874683	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4915 as residues: Gly-22 to Thr-28, Glu-43 to Val-48, Ser-64 to
	Leu-71, Phe-106 to Val-111.
874688	
0/4000	Preferred epitopes include those comprising a sequence shown in SEQ
074600	ID NO. 4917 as residues: Ser-10 to Glu-18, Leu-45 to Arg-54.
874689	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4918 as residues: Asn-13 to Gln-19, Lys-56 to Phe-61, Leu-83
L	to Ala-90.

874695	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4919 as residues: Leu-2 to Ser-12, Pro-125 to Asp-133.
874696	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4920 as residues: Asn-58 to Ser-66.
874699	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4922 as residues: Glu-1 to Ser-7.
874700	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4923 as residues: Gly-10 to Ile-16, Ile-50 to Ser-55.
874701	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4924 as residues: Asn-9 to Gly-14, Glu-17 to His-22.
874702	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4925 as residues: Pro-3 to Arg-20, Pro-24 to Arg-34.
874703	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4926 as residues: Ser-1 to Ser-7, His-35 to Gln-48, Ser-54 to
	Asn-59, Lys-69 to Met-74.
074700	
874708	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4929 as residues: Ala-145 to Gly-152, Val-177 to Gly-185.
874709	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4930 as residues: Ala-13 to Lys-22, Glu-31 to Arg-49, Ser-59 to
	Asn-65.
874710	Preferred epitopes include those comprising a sequence shown in SEQ
6/4/10	
	ID NO. 4931 as residues: Glu-1 to Arg-7, Leu-23 to Arg-39, Lys-46 to
	Asn-52, Pro-59 to Ser-67.
874711	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4932 as residues: Ile-37 to Ala-45, Glu-56 to Pro-62.
874713	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4933 as residues: His-47 to Gly-53, Ser-163 to Ser-169, Pro-
	276 to Lys-282.
974714	Preferred epitopes include those comprising a sequence shown in SEQ
874714	
	ID NO. 4934 as residues: Ser-10 to Glu-18.
874715	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4935 as residues: Ser-13 to Leu-18.
874718	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4937 as residues: Gly-43 to His-54, Phe-126 to Cys-132, Pro-
	140 to Gln-150, Lys-159 to Ala-164, Ser-187 to Gly-193, Pro-212 to
	Gly-227.
074710	
874719	Preferred epitopes include those comprising a sequence shown in SEQ
1	ID NO. 4938 as residues: Gly-1 to Pro-7, Asp-45 to Asp-50, Lys-82 to
1	Leu-89, Asp-97 to His-102, Thr-118 to Ser-126, Phe-128 to Asp-136,
1	Gly-142 to His-148, Ser-212 to Gln-217, Arg-237 to Glu-244, Arg-269
	to Glu-276, Asp-279 to Tyr-284.
874720	Preferred epitopes include those comprising a sequence shown in SEQ
3,4,20	ID NO. 4939 as residues: Glu-18 to Leu-28, Gly-49 to Gly-56, Ser-68 to
	Arg-74.
874724	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4941 as residues: Asp-7 to Glu-12.
874726	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4943 as residues: Ser-55 to Phe-60.
874732	Preferred epitopes include those comprising a sequence shown in SEQ

	ID NO. 4946 as residues: Val-10 to Gly-15, Ser-98 to Thr-105.
874737	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4947 as residues: Ala-36 to His-45.
874741	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4948 as residues: Gln-11 to His-19, Val-30 to Ile-36, Pro-63 to
	Ser-69, Gly-78 to Ser-83, Ser-92 to Tyr-97, Gln-155 to Glu-161, Gly-
	237 to Thr-244.
874744	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4949 as residues: Glu-1 to Phe-12, Ser-47 to Gly-52.
874746	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4951 as residues: Asn-34 to Ser-39.
874749	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4954 as residues: Asp-1 to Gly-17.
874750	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4955 as residues: Gly-4 to Lys-9.
874751	Preferred epitopes include those comprising a sequence shown in SEQ
-	ID NO. 4956 as residues: His-42 to Glu-47.
874752	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4957 as residues: Ile-11 to Gly-17, Gln-26 to Val-32, Gln-41 to
	Asp-52.
874756	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4961 as residues: Ser-1 to His-6.
874757	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4962 as residues: Thr-33 to Phe-38.
874760	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4965 as residues: Gly-1 to Ser-8, Ser-23 to Asn-37.
874763	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4966 as residues: Trp-33 to Gln-40, Cys-64 to Ala-70, Ser-148
	to Tyr-160.
874764	Preferred epitopes include those comprising a sequence shown in SEQ
<u></u>	ID NO. 4967 as residues: Lys-1 to Gln-19.
874765	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4968 as residues: Thr-50 to Gln-59, Ser-62 to Lys-68.
874766	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4969 as residues: Pro-1 to Gly-21, Leu-37 to Pro-42.
874767	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4970 as residues: Lys-30 to Ala-41, Pro-50 to Asn-56, Glu-141
	to Pro-151, Ser-175 to Ser-189.
874769	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4972 as residues: Lys-13 to Glu-22, Glu-76 to Trp-89, Thr-112
	to Gly-120, Arg-141 to Gly-146, Thr-178 to Val-185, Val-212 to Arg-
	223, Pro-225 to Gln-231, Asn-238 to Ala-244, Pro-281 to Glu-287.
874772	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4973 as residues: Gln-44 to Arg-55, Pro-61 to Ala-66.
874774	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4975 as residues: Pro-19 to Pro-34, Leu-46 to Phe-62.
874776	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4977 as residues: Pro-7 to Cys-15, Arg-31 to Glu-42, Ala-47 to
	Ser-58.

874778	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4978 as residues: Arg-1 to Gly-6.
874779	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4979 as residues: Ser-23 to Glu-31, Asp-46 to Pro-53.
874783	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4982 as residues: Gly-1 to Asp-12, Gly-29 to Gly-37, Gly-73 to
	Lys-99.
874784	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4983 as residues: Pro-12 to Gly-18.
874785	Preferred epitopes include those comprising a sequence shown in SEQ
874787	ID NO. 4984 as residues: Lys-24 to Lys-36. Preferred epitopes include those comprising a sequence shown in SEQ
0/4/0/	ID NO. 4986 as residues: Thr-5 to Gly-11, Arg-63 to Lys-73, Gln-92 to
	Glu-98, Ala-106 to Gly-112.
874788	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4987 as residues: Pro-53 to Asn-59.
874790	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4988 as residues: Ser-4 to Thr-9, Gly-17 to Pro-22, Gly-32 to
	Pro-37.
874791	Preferred epitopes include those comprising a sequence shown in SEQ
074702	ID NO. 4989 as residues: Gly-1 to Ser-6, Pro-20 to Arg-27.
874793	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 4990 as residues: Pro-6 to Ala-12, Pro-18 to Thr-28, Pro-31 to
	Arg-37, Pro-53 to Ile-60.
874795	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4991 as residues: Pro-58 to Leu-72.
874796	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4992 as residues: Thr-4 to Arg-11, Pro-30 to Gly-43, Glu-48 to
	Glu-56, Met-86 to Ser-92.
874797	Preferred epitopes include those comprising a sequence shown in SEQ
874800	ID NO. 4993 as residues: Gly-52 to Thr-60, Arg-94 to Glu-100. Preferred epitopes include those comprising a sequence shown in SEQ
8/4800	ID NO. 4994 as residues: Thr-14 to Tyr-25.
874802	Preferred epitopes include those comprising a sequence shown in SEQ
0	ID NO. 4996 as residues: Lys-17 to Leu-23.
874803	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 4997 as residues: Glu-7 to Arg-15, Pro-23 to Arg-36, Pro-79 to
	Ser-96, Ser-119 to Gly-125.
874813	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5004 as residues: Arg-18 to Arg-23, Glu-35 to Asp-50, Ser-67
974915	to Gln-74, Asp-78 to Ser-93.
874815	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5005 as residues: His-38 to Val-46, Ser-97 to Ser-103, Ser-106
1	to Leu-111, His-191 to Gly-196, Leu-223 to Gly-239, Pro-245 to Ala-
	250.
874818	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5007 as residues: Tyr-46 to Gly-51.
874819	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5008 as residues: Pro-33 to Gly-40.

074000	
874820	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5009 as residues: Ile-18 to Gly-30, Leu-33 to Asn-48.
874821	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5010 as residues: Thr-8 to Ser-16.
874822	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5011 as residues: Asn-9 to Phe-14, Glu-63 to Thr-68.
874827	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5012 as residues: Pro-19 to Ser-24, Val-28 to Glu-34.
874828	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5013 as residues: Lys-17 to Gly-28, Thr-62 to Thr-69, Val-88 to
	Arg-101, Gln-106 to Pro-112, Arg-127 to Cys-132, Gly-158 to Leu-163.
874830	Preferred epitopes include those comprising a sequence shown in SEQ
071005	ID NO. 5015 as residues: Arg-53 to Thr-58.
874835	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5017 as residues: Gly-1 to Ser-11, Ser-16 to Ala-26, Thr-28 to
074026	Ser-36, Gln-53 to Trp-59, Lys-72 to Thr-100, Asp-137 to Cys-143.
874836	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5018 as residues: Leu-12 to Asn-17, Phe-25 to Cys-33, Gln-50
874837	to Ser-60, Glu-63 to Pro-68, Pro-83 to Pro-95.
0/403/	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5019 as residues: Val-35 to Thr-41.
874844	Preferred epitopes include those comprising a sequence shown in SEQ
074044	ID NO. 5021 as residues: Pro-19 to Phe-26, Pro-29 to Gly-34, Pro-50 to
	Ser-55, Gly-67 to Lys-73.
874845	Preferred epitopes include those comprising a sequence shown in SEQ
074043	ID NO. 5022 as residues: Asn-1 to Leu-6, Phe-14 to Gly-20.
874847	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5023 as residues: Lys-16 to Thr-22, Glu-36 to Arg-42.
874851	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5025 as residues: Asp-58 to Gly-65, Asp-132 to Cys-147, Pro-
	149 to Pro-157, Pro-218 to Leu-224.
874852	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5026 as residues: Ala-16 to Trp-21.
874854	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5027 as residues: Gly-2 to Glu-8, Met-21 to Trp-26.
874856	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5029 as residues: His-15 to Asp-20, Lys-27 to Asn-33.
874857	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5030 as residues: Lys-35 to Arg-44, Lys-53 to Val-64, Glu-76
	to Val-82, Leu-109 to Lys-118.
874864	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5033 as residues: Leu-40 to Cys-51, Glu-80 to Thr-89, Pro-124
	to Ser-132, Cys-153 to Cys-160, Glu-203 to Asp-209, Ala-226 to Arg-
074065	241.
874865	Preferred epitopes include those comprising a sequence shown in SEQ
074021	ID NO. 5034 as residues: His-1 to Lys-7.
874871	Preferred epitopes include those comprising a sequence shown in SEQ
]	ID NO. 5038 as residues: Gly-1 to Ser-10, Ser-13 to Ile-19, Arg-30 to
	Leu-37, Pro-39 to Asp-48, Pro-140 to Cys-148, Gln-154 to Cys-162,

	Pro-164 to Ser-170.
874873	Preferred epitopes include those comprising a sequence shown in SEQ
0,40,5	ID NO. 5039 as residues: Cys-6 to Ala-12, Pro-14 to Pro-22, Arg-48 to
	Arg-53, Ile-75 to Thr-85, Glu-97 to Gln-102, Arg-130 to Arg-135, Ser-
	147 to Val-152, Lys-175 to Thr-185, Phe-189 to Met-194, Gly-213 to
	Ser-220, Glu-262 to Leu-268.
874879	Preferred epitopes include those comprising a sequence shown in SEQ
.071075	ID NO. 5043 as residues: Glu-1 to Gly-15, His-27 to Thr-39, Gly-43 to
	Ile-49.
874880	Preferred epitopes include those comprising a sequence shown in SEQ
074000	ID NO. 5044 as residues: Pro-62 to Val-70, Lys-103 to Ile-108.
874881	Preferred epitopes include those comprising a sequence shown in SEQ
074001	ID NO. 5045 as residues: Asp-1 to Gly-9.
874885	Preferred epitopes include those comprising a sequence shown in SEQ
074883	ID NO. 5046 as residues: Lys-49 to Gln-55, Glu-83 to Lys-90, Gly-158
	to Gly-164, Lys-185 to Gly-192.
874886	Preferred epitopes include those comprising a sequence shown in SEQ
074000	ID NO. 5047 as residues: Pro-10 to Gly-16, His-128 to Gly-134, His-
	154 to Asp-160, Leu-182 to Leu-187.
874888	Preferred epitopes include those comprising a sequence shown in SEQ
074000	ID NO. 5048 as residues: Pro-15 to Met-27, Thr-106 to His-118, Arg-
	128 to Arg-139, Val-248 to Arg-254.
874889	Preferred epitopes include those comprising a sequence shown in SEQ
0,400	ID NO. 5049 as residues: Pro-7 to Ile-14, Ser-17 to Gln-22.
874890	Preferred epitopes include those comprising a sequence shown in SEQ
0,10,0	ID NO. 5050 as residues: Gly-25 to Ser-31, Trp-34 to Cys-41.
874891	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5051 as residues: Glu-26 to Ser-33, Thr-82 to Phe-90, Met-107
	to Asn-114, Thr-125 to Glu-131, His-175 to Asp-180.
874892	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5052 as residues: Arg-1 to Lys-29, Ile-36 to Lys-47, Lys-52 to
	Gly-83, Pro-89 to Asp-111.
874893	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5053 as residues: Arg-17 to Ile-22.
874896	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5056 as residues: Arg-21 to Lys-26, Pro-37 to Cys-45.
874897	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5057 as residues: Asn-13 to Ala-27, Pro-33 to Lys-42, Asp-61
	to Ser-74, Leu-85 to Lys-102.
874898	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5058 as residues: Pro-1 to Leu-9.
874900	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5060 as residues: Lys-3 to Asp-12, Gln-36 to Tyr-47.
874903	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5062 as residues: Pro-9 to Trp-21, Lys-54 to Gln-61, Lys-75 to
	Phe-87, Glu-97 to Pro-104, Leu-200 to Val-205, Pro-208 to Gly-218,
	Thr-263 to Leu-278.
874905	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5064 as residues: Tyr-94 to Ile-99.

874906	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5065 as residues: Glu-4 to Pro-11.
874907	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5066 as residues: Gln-1 to Lys-10, Thr-17 to Asn-32, Lys-54 to
1	Lys-65.
874908	Preferred epitopes include those comprising a sequence shown in SEQ
4	ID NO. 5067 as residues: Ile-1 to Leu-6, Leu-17 to Ala-23, Ile-27 to
	Thr-33, Asn-40 to Leu-45.
874909	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5068 as residues: Pro-18 to Ser-28, Ser-55 to Thr-64, Asn-90 to
	Lys-95, Asn-128 to Ile-159, Pro-171 to Gly-178, Pro-186 to Lys-192.
874917	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5071 as residues: Arg-37 to Thr-42, Pro-50 to Gly-68, Pro-70 to
	Leu-78, Lys-84 to Lys-89, Asn-95 to Val-105, Asp-117 to Lys-126.
874924	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5072 as residues: Leu-8 to Asn-18, Gly-31 to Ala-39.
874925	Preferred epitopes include those comprising a sequence shown in SEQ
L	ID NO. 5073 as residues: Ser-3 to Arg-9, Gln-24 to Gly-29.
874926	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5074 as residues: Gly-1 to Pro-22.
874928	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5076 as residues: Pro-15 to Gly-23, Ser-27 to Lys-33, Glu-41 to
	Lys-46, Pro-48 to Asp-55.
874937	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5084 as residues: Ser-15 to Ser-20.
874938	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5085 as residues: Ser-12 to Asp-18, His-43 to Gly-51.
874939	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5086 as residues: Ser-12 to Gln-21.
874946	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5088 as residues: Ser-1 to Lys-6, Lys-16 to Glu-24, Asn-34 to
	Lys-47.
874957	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5090 as residues: Ala-12 to Asn-20, Pro-23 to Asn-28, Phe-47
074050	to Val-52, Lys-88 to Gly-93, Tyr-113 to Asn-123, Val-211 to Lys-216.
874958	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5091 as residues: Cys-2 to Leu-9, Pro-37 to Gly-42, Ala-50 to
ļ	Gly-71, Asn-83 to Ala-94, Leu-109 to Leu-115, Phe-156 to Gly-164,
	Lys-234 to His-249, Glu-267 to Gly-281, Asn-335 to Asp-356, Glu-378
	to Ser-385, Gln-402 to Gly-411, Trp-469 to Lys-477, Glu-481 to Gly-486.
874962	
0/4902	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5092 as residues: Asp-1 to Ser-11, Ser-29 to Ser-37, Gln-100 to
1	Arg-112, Leu-123 to Trp-148, Lys-237 to Glu-242, Ala-261 to Asp-266,
	Asp-279 to Ser-300, Thr-374 to Glu-384, Thr-426 to Thr-432, Glu-443 to Val-449.
874965	
5,4,05	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5093 as residues: Asn-13 to His-23, Ser-43 to Gln-56, Val-60 to Glu-65, Pro-67 to Gly-103, Asn-105 to Asp-110.
874970	
017710	Preferred epitopes include those comprising a sequence shown in SEQ

	ID NO. 5094 as residues: Pro-3 to Lys-17, Thr-37 to Gly-47.
874972	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5095 as residues: Thr-124 to Thr-129, Gly-136 to Phe-142,
	Asp-164 to His-171, Asp-180 to Tyr-194.
874973	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5096 as residues: Trp-48 to Arg-56, Pro-68 to Ala-74.
874974	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5097 as residues: Arg-1 to Gly-6, Pro-14 to Ala-26, Ala-42 to
	Lys-47, Pro-66 to Val-82.
874975	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5098 as residues: Ala-18 to Glu-24, Gln-26 to Gln-31.
874976	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5099 as residues: Lys-13 to Ser-19, Pro-33 to Gly-41.
874981	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5104 as residues: Arg-11 to Arg-20.
874983	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5105 as residues: Lys-1 to Thr-9, Ala-43 to Asp-49, Asp-66 to
	Arg-72, Gln-80 to Asp-87, Arg-97 to Lys-104, Ser-111 to Glu-117, Phe-
	150 to Phe-155, Phe-165 to Ala-177, Tyr-219 to Asn-224, Gln-235 to
	Thr-242, Tyr-244 to Thr-251, Arg-267 to Thr-276, Thr-299 to Ile-306,
	Pro-318 to Glu-348, Gly-352 to Leu-370.
874984	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5106 as residues: Thr-40 to Glu-46, Lys-51 to Asn-63.
874991	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5110 as residues: Ser-34 to Gln-40, Met-43 to Asp-70.
874993	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5112 as residues: Thr-6 to Gly-12.
874994	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5113 as residues: Val-3 to Lys-9.
874995	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5114 as residues: Arg-1 to Glu-6, Pro-21 to Thr-27, Lys-41 to
	Thr-48, Gly-202 to Ile-208, Glu-216 to Lys-221, Glu-241 to Lys-247,
	Glu-261 to Leu-267, Pro-269 to Glu-277, Gln-319 to Lys-326.
874996	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5115 as residues: Glu-1 to Gly-12, Tyr-15 to Pro-22, Asp-36 to
	Thr-48.
874997	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5116 as residues: Ile-3 to Lys-9, Ser-31 to Trp-40.
874999	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5118 as residues: Lys-11 to Gln-16.
875002	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5120 as residues: Lys-6 to His-16.
875004	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5122 as residues: Pro-5 to Val-14, Asn-24 to Tyr-35, Ser-70 to
	Val-77, Ser-81 to Asp-99, Ser-121 to Phe-127, Thr-137 to Lys-146, Lys-
	158 to Ser-164, Phe-185 to Gly-192, Asp-212 to Gln-221.
875005	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5123 as residues: Glu-1 to Ser-14.
875008	Preferred epitopes include those comprising a sequence shown in SEQ

	ID NO. 5125 as residues: Arg-1 to Glu-6, Val-14 to Asp-21.
875009	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5126 as residues: Val-30 to Arg-37, Glu-57 to Thr-63, Leu-66
	to Arg-72.
875017	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5128 as residues: Ser-28 to Leu-34, Glu-55 to Gln-62.
875024	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5132 as residues: Tyr-19 to Tyr-24.
875027	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5134 as residues: Thr-46 to Gly-51.
875029	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5135 as residues: Ser-23 to Gly-35.
875034	Preferred epitopes include those comprising a sequence shown in SEQ
1	ID NO. 5137 as residues: Ser-42 to Trp-53, Glu-71 to Ala-78.
875036	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5139 as residues: Ile-20 to Gly-40.
875037	Preferred epitopes include those comprising a sequence shown in SEQ
L	ID NO. 5140 as residues: Trp-23 to Gly-28.
875044	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5143 as residues: Gln-23 to Cys-42, Arg-66 to Asn-73.
875045	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5144 as residues: Glu-10 to Leu-25, Lys-27 to Cys-57.
875046	Preferred epitopes include those comprising a sequence shown in SEQ
L	ID NO. 5145 as residues: Phe-14 to Phe-19.
875049	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5148 as residues: Thr-5 to Lys-12.
875053	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5149 as residues: Ser-16 to Phe-31.
875056	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5151 as residues: Pro-14 to Trp-19.
875058	Preferred epitopes include those comprising a sequence shown in SEO
1	ID NO. 5152 as residues: Pro-3 to Gly-20, Gly-24 to Thr-29, Arg-46 to
	Asn-57, Leu-72 to Phe-78, Glu-81 to Gln-86, Ile-103 to Gln-117, Leu-
	127 to Ile-142, Asn-144 to Ser-151, Arg-156 to His-166.
875060	Preferred epitopes include those comprising a sequence shown in SEO
	ID NO. 5154 as residues: Pro-14 to Ser-20, Pro-41 to Arg-46, Asp-70 to
	His-78.
875062	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5156 as residues: Cys-10 to Tyr-16.
875063	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5157 as residues: Ala-18 to Pro-28.
875066	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5158 as residues: Glu-144 to Leu-152, Glu-170 to Asp-179,
	Gln-225 to Asp-239, Gly-259 to Ala-265.
875067	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5159 as residues: Arg-7 to Pro-16, Pro-37 to Ile-44, Thr-50 to
	Tyr-72, Pro-88 to Phe-94, Ala-107 to Pro-115.
875068	Preferred epitopes include those comprising a sequence shown in SEO
	ID NO. 5160 as residues: Thr-12 to Trp-23.

875070	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5161 as residues: Asp-17 to Asp-27, Pro-34 to Tyr-40, Glu-52
	to Glu-57.
875080	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5163 as residues: Val-30 to Met-37, Glu-39 to Gly-45.
875088	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5165 as residues: Thr-1 to Tyr-8, Gln-27 to Glu-33, Gly-42 to
	Ser-49, Arg-56 to Lys-81, Cys-97 to Lys-104, His-114 to Ser-133, Gln-
	139 to Lys-146, Arg-165 to Glu-173, Asp-180 to Lys-188, Arg-196 to Glu-201.
875092	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5166 as residues: Thr-9 to Asp-17, Leu-70 to Lys-95, Asp-115
	to Leu-124.
875093	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5167 as residues: Gly-2 to Gly-7, Glu-9 to Gln-16, Cys-24 to
	Gly-30, Ala-35 to Ala-45, Ala-55 to Ala-60, Cys-79 to Leu-90, Asp-95
	to Asp-103.
875094	Preferred epitopes include those comprising a sequence shown in SEQ
075100	ID NO. 5168 as residues: His-80 to Glu-87.
875100	Preferred epitopes include those comprising a sequence shown in SEQ
875102	ID NO. 5170 as residues: Thr-18 to Glu-23. Preferred epitopes include those comprising a sequence shown in SEQ
8/3102	ID NO. 5172 as residues: Ser-10 to Gly-16, Pro-24 to Arg-35, Lys-39 to
	Ala-51.
875103	Preferred epitopes include those comprising a sequence shown in SEQ
873103	ID NO. 5173 as residues: Arg-35 to Ala-41.
875105	Preferred epitopes include those comprising a sequence shown in SEQ
375105	ID NO. 5174 as residues: Phe-70 to His-75.
875106	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5175 as residues: His-45 to Gly-55.
875113	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5177 as residues: Thr-27 to Thr-53.
875114	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5178 as residues: Gly-2 to Arg-7.
875118	Preferred epitopes include those comprising a sequence shown in SEQ
Ì	ID NO. 5180 as residues: Pro-21 to Leu-26, Val-62 to Phe-70, Pro-81 to
075101	Asp-89.
875121	Preferred epitopes include those comprising a sequence shown in SEQ
875123	ID NO. 5181 as residues: Phe-19 to Leu-36, Glu-38 to Pro-45. Preferred epitopes include those comprising a sequence shown in SEQ
0/3123	ID NO. 5182 as residues: Ser-44 to Pro-49, Arg-54 to Gly-64, Leu-94 to
	Asp-100, Ser-107 to Gly-113, Lys-143 to Tyr-150.
875126	Preferred epitopes include those comprising a sequence shown in SEQ
0,3120	ID NO. 5185 as residues: His-22 to Ser-27, Cys-34 to Ser-40.
875133	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5187 as residues: His-1 to Gly-9, Gly-19 to Pro-28, Pro-36 to
	Tyr-42, Gly-44 to Gly-65.
875134	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5188 as residues: Gly-10 to Lys-19, Met-21 to Pro-32.

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875143	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5190 as residues: Arg-17 to Ser-23.
875144	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5191 as residues: Asn-14 to Thr-19.
875151	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5193 as residues: Arg-10 to Trp-15, Lys-90 to Ile-95, Asn-103 to Ile-109, Asn-131 to Leu-137, Asn-153 to Arg-163.
875160	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5197 as residues: Val-20 to Asn-27.
875165	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5198 as residues: Thr-5 to Gly-13, Cys-24 to Lys-33.
875177	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5200 as residues: Ala-37 to Asp-44.
875182	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5202 as residues: Pro-25 to Ser-33, Gln-113 to Ser-122, Trp-147 to Tyr-158, Ser-187 to Ala-198, His-201 to Gly-209, Pro-223 to Gly-228, Glu-233 to Gly-238.
875194	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5205 as residues: Ser-16 to Ser-21, Gln-34 to Thr-41.
875200	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5208 as residues: Gln-12 to Cys-19.
875203	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5209 as residues: Arg-1 to Trp-6, Pro-9 to Leu-14.
875205	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5210 as residues: Leu-22 to Ala-27, Ser-31 to Ser-36, Pro-77 to Cys-83.
875206	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5211 as residues: Pro-69 to Pro-75.
875208	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5212 as residues: Asn-25 to Gly-30, Asn-34 to Asn-39.
875209	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5213 as residues: Asn-11 to Ser-18, His-20 to Arg-26, Val-31 to Trp-41.
875210	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5214 as residues: Leu-37 to Thr-52.
875214	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5216 as residues: Ala-7 to Leu-33.
875215	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5217 as residues: Gln-18 to Leu-29, Asp-52 to Ile-57.
875223	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5218 as residues: Thr-2 to Gln-7.
875226	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5219 as residues: Arg-1 to Gln-7, Lys-21 to Gln-31, Leu-41 to Ser-84, Asp-87 to Arg-98, Leu-102 to Lys-115, Leu-129 to Lys-139.
875228	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5220 as residues: Ser-1 to His-10, Pro-84 to Arg-98, His-108 to Asn-113.
875240	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5224 as residues: Ser-31 to Arg-43.

Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5225 as residues: Phe-29 to Leu-37.		
Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5228 as residues: Ser-10 to Asp-24. B75270 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5230 as residues: Ser-1 to Ser-11, Gln-64 to Gln-69, Arg-117 to Pro-128, Pro-135 to Asp-140, Gly-147 to Arg-160, Lys-168 to Val-173, Asn-181 to Lys-191, Glu-200 to Gly-205, Gly-215 to Lys-224. B75271 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5231 as residues: Phe-12 to Lys-17. B75275 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5234 as residues: Pro-9 to Gly-20. B75277 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5234 as residues: Arg-6 to Ser-18. B75278 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5235 as residues: Thr-45 to Lys-50. B75282 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5235 as residues: Thr-45 to Lys-31. B75287 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5240 as residues: Lys-15 to Try-31, Val-44 to Cys-51. B75288 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5240 as residues: Lys-15 to Try-31, Val-44 to Cys-51. B75280 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5241 as residues: Glu-26 to Ala-32, Thr-81 to Ser-50, Arg-61 to Arg-70, Gln-75 to Gly-86. B75296 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5243 as residues: Glu-26 to Ala-32, Thr-81 to Ser-90. B75303 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5243 as residues: Glu-26 to Ala-32, Thr-81 to Ser-90. B75306 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5254 as residues: Pro-36 to Pro-41. B75307 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5254 as residues: Pro-36 to Pro-41. B75310 Preferred epitopes include those comprising a sequence	875246	
ID NO. 5228 as residues: Ser-10 to Asp-24.	875261	
Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5230 as residues: Ser1 to Ser11, Gln-64 to Gln-69, Arg117 to Pro-128, Pro-135 to Asp-140, Gly-147 to Arg160, Lys-168 to Val-173, Asn-181 to Lys-191, Glu-200 to Gly-205, Gly-215 to Lys-224. 875271 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5231 as residues: Phe-12 to Lys-17. 875275 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5232 as residues: Pro-9 to Gly-20. 875277 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5234 as residues: Arg-6 to Ser-18. 875278 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5235 as residues: Thr-45 to Lys-50. 875282 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5239 as residues: Thr-45 to Lys-50. 875287 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5239 as residues: Thr-14 to Lys-31. 875287 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5240 as residues: Lys-15 to Trp-31, Val-44 to Cys-51. 875288 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5241 as residues: Pro-28 to Gly-39, Ser-42 to Ser-50, Arg-61 to Arg-70, Gln-75 to Gly-86. 875296 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5243 as residues: Glu-26 to Ala-32, Thr-81 to Ser-90. 875303 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5243 as residues: Glu-26 to Met-9, Asp-17 to Asn-22, Leu-27 to Val-35. 875306 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5243 as residues: Thr-17 to Phe-22. 875307 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5249 as residues: Pro-36 to Pro-41. 875309 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5250 as residues: Pro-1 to Tyr-22. 875310 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5250 as residues: Ser.		
D NO. 5230 as residues: Ser-1 to Ser-11, Gln-64 to Gln-69, Arg-117 to Pro-128, Pro-135 to Asp-140, Gly-147 to Arg-160, Lys-168 to Val-173, Asn-181 to Lys-191, Glu-200 to Gly-205, Gly-215 to Lys-224. 875271 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5231 as residues: Phe-12 to Lys-17. 875275 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5232 as residues: Pro-9 to Gly-20. 875277 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5234 as residues: Arg-6 to Ser-18. 875278 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5234 as residues: Thr-45 to Lys-50. 875282 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5239 as residues: Thr-14 to Lys-31. 875287 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5240 as residues: Lys-15 to Trp-31, Val-44 to Cys-51. 875288 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5240 as residues: Pro-28 to Gly-39, Ser-42 to Ser-50, Arg-61 to Arg-70, Gln-75 to Gly-86. 875296 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5243 as residues: Glu-26 to Ala-32, Thr-81 to Ser-90. 875303 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5243 as residues: Glu-26 to Ala-32, Thr-81 to Ser-90. 875304 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5243 as residues: Glu-26 to Ala-32, Thr-81 to Ser-90. 875307 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5243 as residues: Thr-17 to Phe-22. 875308 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5248 as residues: Pro-1 to Tyr-22. 875309 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5250 as residues: Fro-1 to Tyr-14, Glu-41 to Leu-49. 875310 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5250 as residues: SPo-1 to Ala-9, Gly-42 to Gln-51. 875311 Preferred epitope	875270	
Pro-128, Pro-135 to Asp-140, Gly-147 to Arg-160, Lys-168 to Val-173, Asn-181 to Lys-191, Glu-200 to Gly-205, Gly-215 to Lys-224. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5231 as residues: Phe-12 to Lys-17. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5232 as residues: Pro-9 to Gly-20. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5234 as residues: Arg-6 to Ser-18. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5235 as residues: Thr-45 to Lys-50. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5239 as residues: Thr-14 to Lys-31. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5240 as residues: Lys-15 to Trp-31, Val-44 to Cys-51. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5240 as residues: Lys-15 to Trp-31, Val-44 to Cys-51. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5241 as residues: Pro-28 to Gly-39, Ser-42 to Ser-50, Arg-61 to Arg-70, Gln-75 to Gly-86. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5243 as residues: Glu-26 to Ala-32, Thr-81 to Ser-90. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5243 as residues: Glu-26 to Ala-32, Thr-81 to Ser-90. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5248 as residues: Pro-11 to Tyr-22. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5249 as residues: Pro-11 to Tyr-22. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5249 as residues: Pro-1 to Tyr-14, Glu-41 to Leu-49. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5250 as residues: Pro-1 to Ala-9, Gly-42 to Gln-51. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5253 as residues: Pro-1 to Ala-9, Gly-42 to Gln-51. Preferred epitopes include those comprising a sequence shown in		
Asn-181 to Lys-191, Glu-200 to Gly-205, Gly-215 to Lys-224. 875271 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5231 as residues: Phe-12 to Lys-17. 875275 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5232 as residues: Pro-9 to Gly-20. 875277 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5234 as residues: Arg-6 to Ser-18. 875278 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5235 as residues: Thr-45 to Lys-50. 875282 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5239 as residues: Thr-14 to Lys-31. 875287 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5240 as residues: Lys-15 to Trp-31, Val-44 to Cys-51. 875288 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5240 as residues: Lys-15 to Trp-31, Val-44 to Cys-51. 875286 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5241 as residues: Pro-28 to Gly-39, Ser-42 to Ser-50, Arg-61 to Arg-70, Gln-75 to Gly-86. 875296 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5243 as residues: Glu-26 to Ala-32, Thr-81 to Ser-90. 875303 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5244 as residues: Glu-26 to Ala-32, Thr-81 to Ser-90. 875306 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5247 as residues: Thr-17 to Phe-22. 875307 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5248 as residues: Pro-1 to Tyr-22. 875308 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5250 as residues: Pro-36 to Pro-41. 875319 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5254 as residues: Pro-36 to Pro-41. 875310 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5254 as residues: Serval to Serval to Serval to Serval to Serval to Serval to Serval to Serval to Serval to Serval		
B75275 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5232 as residues: Pro-9 to Gly-20. 875277 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5234 as residues: Arg-6 to Ser-18. 875278 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5235 as residues: Thr-45 to Lys-50. 875282 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5235 as residues: Thr-14 to Lys-31. 875287 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5240 as residues: Lys-15 to Trp-31, Val-44 to Cys-51. 875288 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5240 as residues: Lys-15 to Trp-31, Val-44 to Cys-51. 875288 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5241 as residues: Pro-28 to Gly-39, Ser-42 to Ser-50, Arg-61 to Arg-70, Gln-75 to Gly-86. 875296 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5243 as residues: Glu-26 to Ala-32, Thr-81 to Ser-90. 875303 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5244 as residues: Glu-26 to Met-9, Asp-17 to Asn-22, Leu-27 to Val-35. 875306 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5247 as residues: Thr-17 to Phe-22. 875307 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5248 as residues: Pro-1 to Tyr-22. 875308 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5250 as residues: Pro-1 to Tyr-24. 875319 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5250 as residues: Pro-1 to Tyr-14, Glu-41 to Leu-49. 875310 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5253 as residues: Gln-23 to Leu-34, Asp-45 to Arg-60. 875311 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5254 as residues: Asp-10 to Lys-16, Lys-35 to Asp-41, Tyr-55 to Leu-62, Glu-145 to Thr-153, Ser-169 to Lys-175, Thr-184 to H		Asn-181 to Lys-191, Glu-200 to Gly-205, Gly-215 to Lys-224.
Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5232 as residues: Pro-9 to Gly-20. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5234 as residues: Arg-6 to Ser-18. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5235 as residues: Thr-45 to Lys-50. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5239 as residues: Thr-14 to Lys-31. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5240 as residues: Lys-15 to Trp-31, Val-44 to Cys-51. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5240 as residues: Pro-28 to Gly-39, Ser-42 to Ser-50, Arg-61 to Arg-70, Gln-75 to Gly-86. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5243 as residues: Glu-26 to Ala-32, Thr-81 to Ser-90. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5243 as residues: Glu-26 to Ala-32, Thr-81 to Ser-90. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5243 as residues: Glu-2 to Met-9, Asp-17 to Asn-22, Leu-27 to Val-35. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5247 as residues: Thr-17 to Phe-22. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5249 as residues: Pro-1 to Tyr-22. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5250 as residues: Pro-1 to Tyr-24. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5253 as residues: Pro-1 to Ala-9, Gly-42 to Gln-51. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5254 as residues: Pro-1 to Ala-9, Gly-42 to Gln-51. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5253 as residues: Pro-15 to Ala-9, Gly-42 to Gln-51. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5254 as residues: Gln-23 to Leu-34, Asp-45 to Arg-60. Preferred epitopes include thos	875271	
ID NO. 5232 as residues: Pro-9 to Gly-20.		
Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5234 as residues: Arg-6 to Ser-18.	875275	
ID NO. 5234 as residues: Arg-6 to Ser-18.	875277	
ID NO. 5235 as residues: Thr-45 to Lys-50.		ID NO. 5234 as residues: Arg-6 to Ser-18.
Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5239 as residues: Thr-14 to Lys-31.	875278	
ID NO. 5239 as residues: Thr-14 to Lys-31. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5240 as residues: Lys-15 to Trp-31, Val-44 to Cys-51. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5241 as residues: Pro-28 to Gly-39, Ser-42 to Ser-50, Arg-61 to Arg-70, Gln-75 to Gly-86. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5243 as residues: Glu-26 to Ala-32, Thr-81 to Ser-90. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5244 as residues: Glu-2 to Met-9, Asp-17 to Asn-22, Leu-27 to Val-35. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5247 as residues: Thr-17 to Phe-22. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5248 as residues: Pro-1 to Tyr-22. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5249 as residues: Pro-1 to Tyr-22. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5250 as residues: Pro-1 to Ala-9, Gly-42 to Gln-51. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5253 as residues: Leu-7 to Tyr-14, Glu-41 to Leu-49. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5254 as residues: Gln-23 to Leu-34, Asp-45 to Arg-60. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5254 as residues: Asn-25 to Tyr-31. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5256 as residues: Asn-25 to Tyr-31. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5256 as residues: Asn-25 to Tyr-31. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5256 as residues: Asn-25 to Tyr-31. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5256 as residues: Asn-25 to Tyr-31. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5256 as residues: Asn-25 to Tyr-31.		
Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5240 as residues: Lys-15 to Trp-31, Val-44 to Cys-51. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5241 as residues: Pro-28 to Gly-39, Ser-42 to Ser-50, Arg-61 to Arg-70, Gln-75 to Gly-86. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5243 as residues: Glu-26 to Ala-32, Thr-81 to Ser-90. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5244 as residues: Glu-2 to Met-9, Asp-17 to Asn-22, Leu-27 to Val-35. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5247 as residues: Thr-17 to Phe-22. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5248 as residues: Pro-1 to Tyr-22. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5249 as residues: Pro-36 to Pro-41. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5250 as residues: Pro-1 to Ala-9, Gly-42 to Gln-51. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5253 as residues: Leu-7 to Tyr-14, Glu-41 to Leu-49. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5254 as residues: Gln-23 to Leu-34, Asp-45 to Arg-60. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5254 as residues: Asn-25 to Tyr-31. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5256 as residues: Asn-25 to Tyr-31. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5256 as residues: Asn-25 to Tyr-31. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5254 as residues: Asn-25 to Tyr-31. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5256 as residues: Asn-25 to Tyr-31.	875282	· · ·
ID NO. 5240 as residues: Lys-15 to Trp-31, Val-44 to Cys-51.		· · · · · · · · · · · · · · · · · · ·
Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5241 as residues: Pro-28 to Gly-39, Ser-42 to Ser-50, Arg-61 to Arg-70, Gln-75 to Gly-86. 875296 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5243 as residues: Glu-26 to Ala-32, Thr-81 to Ser-90. 875303 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5244 as residues: Glu-2 to Met-9, Asp-17 to Asn-22, Leu-27 to Val-35. 875306 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5247 as residues: Thr-17 to Phe-22. 875307 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5248 as residues: Pro-1 to Tyr-22. 875308 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5249 as residues: Pro-36 to Pro-41. 875309 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5250 as residues: Pro-1 to Ala-9, Gly-42 to Gln-51. 875312 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5253 as residues: Leu-7 to Tyr-14, Glu-41 to Leu-49. 875313 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5254 as residues: Gln-23 to Leu-34, Asp-45 to Arg-60. 875316 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5255 as residues: Asn-25 to Tyr-31. 875319 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5256 as residues: Asn-25 to Tyr-31. 875319 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5254 as residues: Asn-25 to Tyr-31. 875319 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5254 as residues: Asn-26 to Tyr-31. 875319 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5254 as residues: Asn-26 to Tyr-31. 875319 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5256 as residues: Asn-26 to Lys-176, Thr-184 to His-192, Gly-224 to Trp-234, Ala-251 to Leu-256, Glu-276 to Asp-281. 875336 Preferred epitopes inclu	875287	· · · · · · · · · · · · · · · · · · ·
ID NO. 5241 as residues: Pro-28 to Gly-39, Ser-42 to Ser-50, Arg-61 to Arg-70, Gln-75 to Gly-86. 875296 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5243 as residues: Glu-26 to Ala-32, Thr-81 to Ser-90. 875303 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5244 as residues: Glu-2 to Met-9, Asp-17 to Asn-22, Leu-27 to Val-35. 875306 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5247 as residues: Thr-17 to Phe-22. 875307 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5248 as residues: Pro-1 to Tyr-22. 875308 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5249 as residues: Pro-36 to Pro-41. 875309 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5250 as residues: Pro-1 to Ala-9, Gly-42 to Gln-51. 875312 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5253 as residues: Leu-7 to Tyr-14, Glu-41 to Leu-49. 875313 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5254 as residues: Gln-23 to Leu-34, Asp-45 to Arg-60. 875316 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5255 as residues: Asp-25 to Tyr-31. 875319 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5256 as residues: Asp-10 to Lys-16, Lys-35 to Asn-41, Tyr-55 to Leu-62, Glu-145 to Thr-153, Ser-169 to Lys-175, Thr-184 to His-192, Gly-224 to Trp-234, Ala-251 to Leu-256, Glu-276 to Asp-281. 875336 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5261 as residues: Tyr-3 to Leu-10.		
Arg-70, Gln-75 to Gly-86. 875296 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5243 as residues: Glu-26 to Ala-32, Thr-81 to Ser-90. 875303 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5244 as residues: Glu-2 to Met-9, Asp-17 to Asn-22, Leu-27 to Val-35. 875306 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5247 as residues: Thr-17 to Phe-22. 875307 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5248 as residues: Pro-1 to Tyr-22. 875308 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5249 as residues: Pro-36 to Pro-41. 875309 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5250 as residues: Pro-1 to Ala-9, Gly-42 to Gln-51. 875312 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5253 as residues: Leu-7 to Tyr-14, Glu-41 to Leu-49. 875313 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5254 as residues: Gln-23 to Leu-34, Asp-45 to Arg-60. 875316 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5255 as residues: Asp-25 to Tyr-31. 875319 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5256 as residues: Asp-10 to Lys-16, Lys-35 to Asn-41, Tyr-55 to Leu-62, Glu-145 to Thr-153, Ser-169 to Lys-175, Thr-184 to His-192, Gly-224 to Trp-234, Ala-251 to Leu-256, Glu-276 to Asp-281. 875336 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5261 as residues: Tyr-3 to Leu-10.	875288	
Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5243 as residues: Glu-26 to Ala-32, Thr-81 to Ser-90. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5244 as residues: Glu-2 to Met-9, Asp-17 to Asn-22, Leu-27 to Val-35. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5247 as residues: Thr-17 to Phe-22. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5248 as residues: Pro-1 to Tyr-22. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5249 as residues: Pro-36 to Pro-41. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5250 as residues: Pro-1 to Ala-9, Gly-42 to Gln-51. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5253 as residues: Leu-7 to Tyr-14, Glu-41 to Leu-49. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5254 as residues: Gln-23 to Leu-34, Asp-45 to Arg-60. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5255 as residues: Asn-25 to Tyr-31. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5256 as residues: Asn-25 to Tyr-31. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5256 as residues: Asn-25 to Tyr-31. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5256 as residues: Asn-10 to Lys-16, Lys-35 to Asn-41, Tyr-55 to Leu-62, Glu-145 to Thr-153, Ser-169 to Lys-175, Thr-184 to His-192, Gly-224 to Trp-234, Ala-251 to Leu-256, Glu-276 to Asp-281. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5261 as residues: Tyr-3 to Leu-10.	1	-
ID NO. 5243 as residues: Glu-26 to Ala-32, Thr-81 to Ser-90. 875303 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5244 as residues: Glu-2 to Met-9, Asp-17 to Asn-22, Leu-27 to Val-35. 875306 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5247 as residues: Thr-17 to Phe-22. 875307 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5248 as residues: Pro-1 to Tyr-22. 875308 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5249 as residues: Pro-36 to Pro-41. 875309 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5250 as residues: Pro-1 to Ala-9, Gly-42 to Gln-51. 875312 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5253 as residues: Leu-7 to Tyr-14, Glu-41 to Leu-49. 875313 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5254 as residues: Gln-23 to Leu-34, Asp-45 to Arg-60. 875316 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5255 as residues: Asn-25 to Tyr-31. 875319 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5256 as residues: Asp-10 to Lys-16, Lys-35 to Asn-41, Tyr-55 to Leu-62, Glu-145 to Thr-153, Ser-169 to Lys-175, Thr-184 to His-192, Gly-224 to Trp-234, Ala-251 to Leu-256, Glu-276 to Asp-281. 875336 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5261 as residues: Tyr-3 to Leu-10.		
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	875336	
875338 Preferred epitopes include those comprising a sequence shown in SEQ		
	875338	Preferred epitopes include those comprising a sequence shown in SEQ

	ID NO. 5262 as residues: Pro-9 to Ile-14, Glu-81 to Gln-90.
875346	
0/3340	Preferred epitopes include those comprising a sequence shown in SEQ
075247	ID NO. 5264 as residues: Gly-29 to Arg-44.
875347	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5265 as residues: Ile-3 to Ser-14, Ala-32 to Ser-44, Ser-60 to
	Leu-67.
875360	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5268 as residues: Pro-14 to Leu-19, Ile-37 to Ala-46, Ser-58 to
	Asn-65, Pro-71 to Gly-77.
875364	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5269 as residues: Val-38 to Phe-47, Asn-64 to Phe-69.
875367	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5271 as residues: Gly-14 to Leu-21, Asn-31 to Met-37.
875371	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5272 as residues: Pro-12 to Glu-23, Lys-29 to Pro-34, Pro-54 to
	Leu-66.
875372	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5273 as residues: Ala-7 to Arg-12.
875373	Preferred epitopes include those comprising a sequence shown in SEQ
0.5575	ID NO. 5274 as residues: Tyr-54 to Cys-61, Asn-73 to Pro-78, Pro-84 to
	Asn-93, Gln-99 to Asp-105.
875378	Preferred epitopes include those comprising a sequence shown in SEQ
0/33/0	ID NO. 5276 as residues: Leu-42 to Lys-53, Cys-100 to Asn-110, Pro-
	137 to Gly-144, Pro-190 to Ala-205.
875379	Dueformed emitteness in slude these commissions are seen at 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1
0/33/9	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5277 as residues: Asp-5 to Ala-10, Ala-19 to Ile-25, Val-39 to
075201	Ser-44, Gln-74 to Cys-90, Leu-94 to Glu-99, Leu-108 to Phe-116.
875381	Preferred epitopes include those comprising a sequence shown in SEQ
075200	ID NO. 5279 as residues: Cys-46 to Leu-51.
875382	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5280 as residues: Pro-11 to Thr-16, Pro-23 to Gly-33, Ala-51 to
077001	Arg-61.
875384	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5281 as residues: Gln-15 to Gly-28, Asp-83 to Tyr-92.
875385	Preferred epitopes include those comprising a sequence shown in SEQ
<u></u>	ID NO. 5282 as residues: Leu-3 to Asp-8, Gln-30 to His-36.
875388	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5283 as residues: Thr-2 to Ser-9, Pro-23 to Arg-30.
875391	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5284 as residues: Lys-1 to Arg-10, Lys-53 to Tyr-62.
875397	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5285 as residues: Arg-7 to Gly-29, Arg-37 to Glu-47, Asp-78 to
	Thr-83, Gly-173 to Val-180, Glu-188 to Glu-202, Pro-208 to Thr-216,
	Thr-227 to Glu-242, Arg-250 to Gly-281, Lys-288 to Thr-296, Glu-301
	to Arg-311, Ala-313 to Lys-318, Lys-357 to Thr-367, Pro-376 to Ser-
	387, Pro-416 to Lys-428, Pro-486 to Thr-491, Ser-497 to Arg-516, Lys-
	522 to Lys-532, Arg-537 to Met-557.
875402	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5286 as residues: Asn-1 to Thr-15.
L	

075406	Preferred epitopes include those comprising a sequence shown in SEQ
875406	ID NO. 5288 as residues: Pro-5 to Ala-19.
875410	Preferred epitopes include those comprising a sequence shown in SEQ
8/3410	ID NO. 5289 as residues: Ala-4 to Pro-14, Pro-23 to Thr-28, Thr-40 to
	Gln-45, Tyr-60 to Gln-69, Pro-88 to Leu-93, Glu-108 to Ala-113, Val-
	119 to Gly-131, Arg-146 to Arg-155, Ala-164 to Lys-171, Thr-190 to
	Met-201.
875415	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5290 as residues: Arg-18 to Trp-23, Gly-25 to Gly-32, Lys-34
	to Arg-42, Gly-52 to Thr-59, Ala-86 to Lys-92.
875416	Preferred epitopes include those comprising a sequence shown in SEQ
673410	ID NO. 5291 as residues: Lys-9 to Gly-37.
075417	
875417	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5292 as residues: Glu-2 to Cys-14.
875419	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5294 as residues: Thr-2 to Tyr-11.
875423	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5295 as residues: Lys-13 to Ile-24, Phe-28 to Val-35.
875428	Preferred epitopes include those comprising a sequence shown in SEQ
075420	ID NO. 5298 as residues: Gly-2 to Thr-7, Gly-20 to Thr-29, Asn-69 to
075100	Arg-77.
875429	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5299 as residues: Phe-4 to Pro-9, Pro-13 to Gln-18.
875433	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5300 as residues: Lys-78 to Met-83.
875434	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5301 as residues: Thr-34 to Glu-39.
875437	Preferred epitopes include those comprising a sequence shown in SEQ
0,343,	ID NO. 5302 as residues: Glu-1 to Gln-7.
875440	Preferred epitopes include those comprising a sequence shown in SEQ
0/3440	ID NO. 5303 as residues: Arg-11 to Met-17, Ile-66 to Trp-71, Asp-91 to
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	Leu-97, Ala-102 to Lys-111, Trp-113 to Glu-120, Pro-132 to Asn-141,
	Thr-144 to Glu-153, Glu-159 to Glu-172, Pro-177 to Lys-192.
875441	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5304 as residues: Cys-28 to Cys-34.
875442	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5305 as residues: Pro-18 to Lys-23.
875446	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5306 as residues: Pro-8 to Phe-18.
875452	Preferred epitopes include those comprising a sequence shown in SEQ
673432	ID NO. 5307 as residues: Ala-6 to Cys-17.
075150	
875458	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5308 as residues: Glu-40 to Glu-46, Arg-51 to Ser-67.
875462	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5311 as residues: Ser-2 to Ser-14, Arg-75 to Asn-85.
875468	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5313 as residues: Thr-35 to Thr-49.
875474	Preferred epitopes include those comprising a sequence shown in SEQ
0/34/4	ID NO. 5314 as residues: Asp-1 to Asp-13, Arg-40 to Arg-56, Ser-72 to
	THO. 3314 as residues. Asp-1 to Asp-13, Arg-40 to Arg-30, Ser-72 to

	Asp-84, Ala-88 to Arg-96, Lys-115 to Phe-121, Asp-133 to Lys-139,
975475	Leu-203 to Leu-210, Asp-264 to Arg-269.
875475	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5315 as residues: Pro-12 to Gly-19.
875479	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5318 as residues: His-32 to Lys-40.
875481	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5319 as residues: Arg-22 to Ser-39, Val-42 to Thr-54, Gln-61 to
	His-69, Glu-83 to Gly-109, Pro-111 to Gly-118.
875490	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5322 as residues: Cys-75 to Thr-81.
875491	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5323 as residues: Gln-8 to His-15, Ser-32 to Gln-43, Leu-51 to
	Glu-70.
875499	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5329 as residues: Asn-36 to Leu-55.
875500	Preferred epitopes include those comprising a sequence shown in SEQ
0,2500	ID NO. 5330 as residues: Thr-31 to Arg-39.
875501	Preferred epitopes include those comprising a sequence shown in SEQ
0,5501	ID NO. 5331 as residues: Asp-52 to Asn-59.
875508	Preferred epitopes include those comprising a sequence shown in SEQ
073300	ID NO. 5334 as residues: Pro-1 to Ile-18, Asp-28 to Lys-33, Leu-50 to
	Gln-55, Glu-85 to Ala-94, Leu-121 to Ser-130, Lys-143 to Gly-150,
	Leu-173 to Asp-179, Lys-183 to Asp-192, Lys-196 to Glu-202, Asn-219
	to Asn-227, Glu-235 to Glu-248.
875512	Preferred epitopes include those comprising a sequence shown in SEQ
0/3312	ID NO. 5335 as residues: Asp-10 to Trp-16, Glu-33 to Asn-43.
875514	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5336 as residues: Asp-11 to Tyr-32, Gln-43 to Thr-58, His-70
	to Arg-79, Ser-101 to Ala-108, Met-110 to Ser-124.
875515	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5337 as residues: Met-1 to Arg-8, Met-10 to His-17.
875516	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5338 as residues: Leu-2 to Ser-8, Gln-41 to Gly-46, Asp-70 to
	Gln-80, Pro-82 to Gly-97.
875518	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5340 as residues: Arg-1 to Trp-11, Ser-28 to Leu-42, Gly-65 to
	Gly-70, Ala-72 to Gln-77, Gly-89 to Lys-98, Asp-126 to Thr-136, Gln-
	218 to Gly-226, Lys-261 to Gly-282.
875520	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5341 as residues: Arg-5 to Ser-18, Arg-36 to Gly-42, Gln-45 to
	Gly-56, Val-69 to Arg-75.
875525	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5343 as residues: Arg-6 to Thr-22, Arg-31 to His-38.
875527	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5344 as residues: Gly-24 to Leu-31, Ser-64 to Val-70, Arg-93 to
	Trp-100.
875528	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5345 as residues: Thr-6 to Ile-13.
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875534	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5347 as residues: Arg-1 to Thr-14, Arg-28 to Asp-34, Gln-51 to
	Ser-60, Lys-69 to Gly-78, Val-110 to Val-115, Asn-135 to Glu-141,
	Asn-167 to Pro-179, Lys-203 to Lys-214, Gly-267 to Pro-279.
875538	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5348 as residues: Thr-1 to Arg-6.
875544	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5351 as residues: Gln-1 to Asn-8.
875545	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5352 as residues: Cys-2 to Gly-16, Glu-35 to Leu-40, Pro-61 to
	Gln-66.
875547	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5354 as residues: Leu-37 to His-43.
875548	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5355 as residues: Val-15 to Asp-21, Cys-29 to Ser-36.
875550	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5356 as residues: Arg-81 to Gln-93, Leu-103 to Val-116.
875551	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5357 as residues: Glu-11 to Lys-22, Glu-36 to Gly-41.
875553	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5359 as residues: Arg-6 to Lys-11, Phe-16 to Ile-21, Thr-48 to
	Leu-56, Pro-64 to Arg-70.
875554	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5360 as residues: Tyr-2 to Ser-10, Asn-69 to Leu-80.
875559	Preferred epitopes include those comprising a sequence shown in SEQ
1	ID NO. 5363 as residues: Pro-123 to Asn-130.
875563	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5365 as residues: Pro-35 to Gly-62.
875565	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5367 as residues: Pro-2 to Asp-7, Gln-13 to Gln-29, Pro-35 to
	Trp-41.
875570	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5369 as residues: Leu-1 to Ser-6, Ser-45 to Lys-56, Asn-66 to
	Lys-78.
875574	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5372 as residues: Pro-10 to Gln-15, Cys-25 to Ile-30, Ser-42 to
	Lys-47.
875583	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5374 as residues: Lys-6 to Lys-37, Arg-43 to Leu-49, Met-53 to
	Val-59.
875590	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5380 as residues: Cys-128 to Pro-134.
875594	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5381 as residues: Gly-40 to Ser-45.
875596	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5382 as residues: Gly-1 to Gly-10.
875597	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5383 as residues: His-3 to Ser-9.
875604	Preferred epitopes include those comprising a sequence shown in SEQ
	1 - 1 - 1 - 1 - 1 - 1 - 1 - 1 - 1 - 1 -

	ID NO. 5386 as residues: Lys-7 to Ser-20, Arg-67 to Ser-74.
875605	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5387 as residues: Gly-17 to Ser-24, Met-42 to Arg-48.
875606	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5388 as residues: Tyr-1 to Gly-13, Glu-32 to Asp-43, Ser-55 to
	Ile-62, Pro-119 to Asn-131.
875609	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5390 as residues: Thr-12 to Ser-20, Leu-60 to Ala-66.
875610	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5391 as residues: Cys-41 to Ser-47.
875613	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5394 as residues: Leu-12 to Lys-18.
875625	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5395 as residues: Asp-8 to Leu-25, Arg-94 to Ala-102, Glu-133
	to Ala-138.
875628	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5396 as residues: Ser-17 to Gly-23.
875629	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5397 as residues: Glu-1 to Glu-11, Arg-21 to Ser-27.
875631	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5399 as residues: Val-37 to Asn-43, Glu-62 to Pro-69, Gln-118
-	to Tyr-131, Ser-144 to Trp-150.
875633	Preferred epitopes include those comprising a sequence shown in SEQ
075624	ID NO. 5401 as residues: Asn-11 to Arg-16.
875634	Preferred epitopes include those comprising a sequence shown in SEQ
975625	ID NO. 5402 as residues: Ile-1 to Gly-10, Asp-24 to Arg-29.
875635	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5403 as residues: Phe-1 to Ile-8, Thr-21 to Leu-38, Glu-55 to
875636	Lys-70, Lys-76 to Leu-82, Lys-84 to Glu-89, Ile-93 to Ser-98.
873030	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5404 as residues: Pro-30 to Asp-35.
875638	Preferred enitones include these community and the second services and the second services and the second services and the second services are services as the second services are services are services are services as the second services are services ar
075050	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5405 as residues: Asp-1 to Gly-7, Arg-13 to Arg-18, Arg-48 to
	Ser-54.
875640	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5407 as residues: Thr-36 to Cys-47.
875642	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5409 as residues: Arg-2 to Thr-8, Thr-46 to His-51.
875646	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5410 as residues: Ala-4 to Arg-10, Cys-22 to Lys-27.
875650	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5411 as residues: Glu-29 to Lys-34, Leu-151 to Tyr-156, Glu-
	162 to Arg-170.
875651	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5412 as residues: Leu-119 to Gln-125, Arg-128 to Ser-139,
	Gln-145 to Pro-158.
875653	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5413 as residues: Pro-1 to Gln-14.
875654	Preferred epitopes include those comprising a sequence shown in SEQ

	ID NO. 5414 as residues: Arg-34 to Gly-66.
875658	Preferred epitopes include those comprising a sequence shown in SEQ
873038	ID NO. 5415 as residues: His-19 to Tyr-30, Ala-53 to Ala-59, Ala-90 to
	Pro-101, Lys-132 to Lys-139, Ala-152 to Arg-158, Phe-168 to Leu-175,
1	Arg-178 to Lys-186.
875661	Preferred epitopes include those comprising a sequence shown in SEQ
8/3001	
975669	ID NO. 5416 as residues: Tyr-2 to Ser-8, Thr-15 to Ala-25.
875662	Preferred epitopes include those comprising a sequence shown in SEQ
075660	ID NO. 5417 as residues: Gly-5 to Cys-12, Phe-40 to Thr-47.
875663	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5418 as residues: Thr-4 to Ser-12.
875665	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5419 as residues: Lys-2 to Lys-7.
875669	Preferred epitopes include those comprising a sequence shown in SEQ
· · · · · · · · · · · · · · · · · · ·	ID NO. 5420 as residues: Lys-1 to Gly-11.
875677	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5423 as residues: Gly-1 to His-7, Val-10 to Phe-17, Asp-62 to
	Arg-67.
875678	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5424 as residues: Ile-2 to Ile-9, Asn-76 to Gln-82.
875681	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5426 as residues: Glu-1 to Asn-12, Pro-20 to Ala-26, Thr-42 to
	Ser-50.
875683	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5428 as residues: Val-60 to Pro-69.
875687	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5429 as residues: Asp-18 to Phe-24.
875688	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5430 as residues: Glu-8 to Glu-13.
875689	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5431 as residues: Lys-24 to Lys-30.
875690	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5432 as residues: Gly-3 to Leu-20, Trp-38 to Arg-44, Lys-58 to
	Lys-64.
875698	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5434 as residues: Tyr-43 to Lys-52, Glu-60 to Arg-66, Gln-84
	to Cys-89, Gln-106 to Lys-117, Thr-140 to Asp-168, Gln-170 to Arg-
	177.
875704	Preferred epitopes include those comprising a sequence shown in SEQ
3.2,0	ID NO. 5438 as residues: Gly-24 to Thr-30, Ser-103 to Gly-109.
875717	Preferred epitopes include those comprising a sequence shown in SEQ
0,5,1,	ID NO. 5441 as residues: Cys-12 to Cys-34, Pro-36 to Thr-45, Arg-75
	to Asn-85.
875719	Preferred epitopes include those comprising a sequence shown in SEQ
0/3/19	ID NO. 5442 as residues: Asn-1 to Tyr-7.
875722	Preferred epitopes include those comprising a sequence shown in SEQ
0/3/22	ID NO. 5443 as residues: Leu-2 to Phe-7.
075724	
875724	Preferred epitopes include those comprising a sequence shown in SEQ
L	ID NO. 5444 as residues: Asn-86 to Ser-91.

875725 Preferred epitopes include those comprising a sequence shown in	SEQ
ID NO. 5445 as residues: Thr-9 to Thr-17, Arg-33 to Val-41.	
875727 Preferred epitopes include those comprising a sequence shown in	SEQ
ID NO. 5446 as residues: Thr-16 to Pro-23, Pro-39 to Trp-48, Ar	g-50 to
Lys-55, Gly-73 to Gly-79.	
875734 Preferred epitopes include those comprising a sequence shown in	SEQ
ID NO. 5451 as residues: Ser-12 to Thr-18, Pro-20 to Pro-25.	
875736 Preferred epitopes include those comprising a sequence shown in	
ID NO. 5452 as residues: Phe-10 to Arg-15, Ile-48 to Thr-53, Ser	-64 to
Asn-69.	
875737 Preferred epitopes include those comprising a sequence shown in	
ID NO. 5453 as residues: Leu-1 to Cys-6, Ala-74 to Gly-87, Gln-	106 to
Gly-111.	
Preferred epitopes include those comprising a sequence shown in	SEQ
ID NO. 5454 as residues: Glu-11 to Asp-19, Gly-40 to Thr-47, Pr to Arg-71.	0-66
875739 Preferred epitopes include those comprising a sequence shown in	SEO
ID NO. 5455 as residues: Gly-45 to Asp-50.	SEQ
875740 Preferred epitopes include those comprising a sequence shown in	SEO
ID NO. 5456 as residues: Glu-1 to Gln-22.	SLQ
875746 Preferred epitopes include those comprising a sequence shown in	SEO
ID NO. 5457 as residues: Leu-55 to Gln-64.	DDQ
875751 Preferred epitopes include those comprising a sequence shown in	SEO
ID NO. 5459 as residues: Phe-21 to Leu-26, Gly-81 to His-87.	
875752 Preferred epitopes include those comprising a sequence shown in	SEQ
ID NO. 5460 as residues: Ser-11 to Asn-16, Trp-33 to Arg-49.	
875753 Preferred epitopes include those comprising a sequence shown in	SEQ
ID NO. 5461 as residues: Glu-1 to Ile-17, Leu-54 to Asn-59.	
875754 Preferred epitopes include those comprising a sequence shown in	SEQ
ID NO. 5462 as residues: Arg-53 to Val-58.	
875760 Preferred epitopes include those comprising a sequence shown in	SEQ
ID NO. 5463 as residues: Phe-45 to Asn-51.	
875765 Preferred epitopes include those comprising a sequence shown in	SEQ
ID NO. 5465 as residues: Pro-7 to Gly-12. 875766 Preferred epitopes include those comprising a sequence shown in	ar.
875766 Preferred epitopes include those comprising a sequence shown in ID NO. 5466 as residues: Gly-21 to Phe-28.	SEQ
	CEO
875769 Preferred epitopes include those comprising a sequence shown in ID NO. 5468 as residues: Lys-7 to Gly-12.	sey
875772 Preferred epitopes include those comprising a sequence shown in	SEC
ID NO. 5469 as residues: Arg-19 to Pro-45, Gly-60 to Leu-72, Le	
to Asn-109.	u-70
875774 Preferred epitopes include those comprising a sequence shown in	SEO
ID NO. 5471 as residues: Ile-27 to Val-33, Val-63 to Ser-68.	~~~
875779 Preferred epitopes include those comprising a sequence shown in	SEO
ID NO. 5473 as residues: Gln-54 to Ser-63, Glu-84 to Lys-92, Va	1-100
to Gln-105.	1
875781 Preferred epitopes include those comprising a sequence shown in	SEQ
ID NO. 5475 as residues: Glu-72 to Ala-80.	
875783 Preferred epitopes include those comprising a sequence shown in	SEQ

	ID NO. 5477 as residues: Gly-1 to Asn-15.
875784	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5478 as residues: Glu-17 to Asp-22, Asn-30 to Cys-35, Leu-39
	to Lys-49.
875786	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5480 as residues: Arg-8 to Thr-17.
875787	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5481 as residues: Ser-3 to Pro-16, Asp-38 to Ser-43, Arg-53 to
	Gln-62, Trp-78 to Ser-84.
875789	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5482 as residues: Arg-1 to Ile-8, Pro-50 to Thr-62.
875794	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5484 as residues: Thr-8 to Val-13, Tyr-39 to Arg-46.
875800	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5486 as residues: Tyr-1 to Gln-12, Gly-17 to Cys-26, Trp-37 to
	Asn-43, Leu-46 to Gly-51.
875804	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5488 as residues: Asp-54 to Gly-67.
875805	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5489 as residues: Ser-1 to Thr-9.
875809	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5491 as residues: Asn-16 to Leu-30, Ala-48 to Thr-53, Arg-109
	to Asp-114, Arg-120 to Gly-126, Pro-153 to Asp-161, Asn-177 to Lys-
	186, Ser-253 to Ser-260.
875810	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5492 as residues: Pro-1 to Lys-11, Pro-31 to Leu-39, Thr-67 to
	Lys-77.
875814	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5493 as residues: His-1 to Gly-14, Ala-21 to Arg-30.
875815	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5494 as residues: Ile-14 to Leu-35, Pro-37 to Thr-51.
875817	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5496 as residues: Ser-15 to Ile-24, Asn-56 to Lys-67, Ser-80 to
	Lys-95, Gly-148 to Pro-165.
875820	Preferred epitopes include those comprising a sequence shown in SEQ
1	ID NO. 5498 as residues: Phe-2 to Ser-9, Cys-12 to Ser-23, Glu-37 to
	Pro-48, Glu-56 to Asp-64.
875821	Preferred epitopes include those comprising a sequence shown in SEQ
075000	ID NO. 5499 as residues: Gly-98 to Ala-110.
875822	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5500 as residues: Ala-7 to Pro-18, Ser-57 to Ser-64, Phe-94 to
075005	Gln-105, Leu-129 to Gly-141.
875825	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5502 as residues: Lys-1 to Lys-19, Glu-66 to Gln-73, Asn-75 to
075000	Asn-80, Met-112 to Asn-118, Val-122 to Thr-134.
875828	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5504 as residues: His-1 to Leu-12, Leu-16 to Cys-30, Thr-46 to
075022	Asn-56.
875832	Preferred epitopes include those comprising a sequence shown in SEQ

	ID NO. 5505 as residues: Lys-1 to Arg-9, Cys-32 to Tyr-39, Lys-53 to Gly-64, Phe-86 to Asn-92.
875836	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5508 as residues: His-79 to Ser-92.
875837	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5509 as residues: Ser-47 to Arg-54.
875838	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5510 as residues: Ser-1 to Phe-8.
875839	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5511 as residues: Gln-1 to Gly-22, Pro-36 to Arg-42, Arg-89 to Gln-94.
875840	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5512 as residues: Thr-6 to Asn-16, Gln-50 to Lys-66.
875841	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5513 as residues: Ala-44 to Arg-51, Val-71 to Ser-76.
875845	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5514 as residues: Gly-1 to Lys-6, Ser-54 to Ser-60.
875846	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5515 as residues: Ser-28 to Gly-33.
875855	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5521 as residues: Glu-13 to Asn-18, Asn-53 to Lys-59.
875856	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5522 as residues: Ala-28 to Ser-33.
875858	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5523 as residues: His-1 to Asn-17, Gly-21 to Arg-28, Lys-43 to Asn-49, Ser-64 to His-80, Ala-91 to Asp-130, Gly-144 to Ser-158.
875863	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5524 as residues: Pro-23 to Asp-28, Pro-40 to Gln-47.
875864	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5525 as residues: Pro-1 to Ser-15, Leu-27 to Lys-32, Arg-39 to Ser-53, Thr-58 to Glu-81, Gly-87 to Leu-92, Val-96 to Glu-106, Lys-114 to Ile-131, Asp-134 to Lys-140, Asn-142 to Lys-149, Lys-155 to Gly-168.
875865	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5526 as residues: His-11 to Cys-23, Ala-29 to Gln-35, His-43 to Arg-50.
875868	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5527 as residues: Arg-33 to Glu-42, Arg-45 to Gly-64, Ala-79 to Asn-117, Thr-130 to Lys-143, Ser-222 to Lys-233, Val-235 to Asn-240, Leu-289 to Met-294.
875871	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5528 as residues: Gln-1 to Ala-17, Gln-43 to Asp-48.
875874	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5529 as residues: Glu-40 to Thr-50.
875884	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5530 as residues: Ser-14 to Cys-19, Lys-53 to Asn-58, Ser-63 to Ser-70, Gly-118 to Cys-123, Cys-132 to Gly-138.
875886	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5531 as residues: Asn-46 to Glu-51.

875888	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5532 as residues: Lys-1 to Gly-17, Arg-56 to Gln-61, Gln-82 to Pro-89.
875891	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5533 as residues: Tyr-4 to Gly-11, Phe-33 to Asn-38.
875894	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5534 as residues: Arg-11 to Glu-24, Arg-39 to Glu-52, His-70 to Gly-82, His-98 to Arg-124, His-126 to Ser-142, His-154 to Gly-166.
875897	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5535 as residues: Pro-1 to Lys-8, Phe-49 to Pro-67, Leu-88 to Trp-100.
875905	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5539 as residues: Pro-19 to Cys-28, Leu-40 to Thr-49, Glu-57 to Pro-69, Phe-82 to Asn-89.
875908	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5542 as residues: Val-27 to Gly-34.
875912	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5543 as residues: Lys-5 to Gln-11, Ser-16 to Lys-28, Pro-39 to Phe-44, Thr-136 to Lys-148, Cys-182 to His-189, Val-197 to Tyr-202, Ser-273 to Gly-300.
875914	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5545 as residues: Ser-7 to Lys-13, Met-16 to Trp-21, Pro-54 to Gly-60, Ser-112 to Gly-117.
875923	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5547 as residues: Asn-1 to Lys-10, Glu-29 to Thr-35, Glu-41 to Glu-57, Glu-78 to Arg-83, Ala-97 to Trp-102.
875924	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5548 as residues: Gln-1 to Asn-8, Arg-22 to Leu-28, Ser-30 to Phe-48, Ser-51 to Glu-56, Gln-70 to Leu-88, Phe-101 to Asn-111, Arg-113 to Tyr-121, Ser-130 to Asn-135, Glu-141 to Gln-152, Asn-169 to Trp-191.
875925	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5549 as residues: Ser-45 to Ala-50.
875926	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5550 as residues: Leu-4 to Ser-13.
875927	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5551 as residues: Arg-2 to Lys-21.
875932	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5552 as residues: Asp-27 to Gln-33.
875933	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5553 as residues: Gly-1 to Gln-8, Met-19 to Ser-24.
875935	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5555 as residues: Asn-20 to Thr-25, Ser-30 to Pro-35.
875936	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5556 as residues: Gly-12 to Lys-18, Arg-46 to Glu-56, Leu-67 to Gly-73, Ala-91 to Ser-102.
875937	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5557 as residues: Arg-4 to Thr-10, Arg-61 to Glu-71, Leu-82 to Gly-88, Ala-106 to Lys-142.

875939	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5559 as residues: Arg-3 to Leu-15, Arg-17 to Asn-24.
875940	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5560 as residues: Gly-28 to Phe-34, Gly-36 to Cys-41, Arg-46 to Arg-54, Pro-75 to Arg-90.
875941	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5561 as residues: Gln-24 to Glu-35, Lys-53 to Gln-67, Pro-85 to Trp-98.
875942	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5562 as residues: Cys-74 to Ala-84.
875946	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5563 as residues: Gly-34 to Pro-48, Arg-86 to Gly-91.
875951	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5565 as residues: Pro-31 to Leu-41.
875955	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5568 as residues: His-19 to Asn-24, Pro-39 to Lys-45.
875967	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5569 as residues: Arg-30 to Arg-38.
875971	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5570 as residues: Ser-1 to Asp-8, Asn-16 to Ser-35, Asn-47 to Pro-70.
875972	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5571 as residues: Pro-14 to Arg-23, Phe-41 to Gly-49, His-69 to His-76, Tyr-84 to Asn-90.
875976	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5573 as residues: Tyr-3 to Gly-10.
875984	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5576 as residues: Ser-2 to Gln-15.
875991	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5579 as residues: Thr-47 to Gly-53.
875995	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5581 as residues: Pro-3 to Glu-8.
875999	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5584 as residues: Gly-11 to Ala-16, Gln-70 to His-78.
876006	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5586 as residues: Pro-12 to Thr-22.
876008	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5588 as residues: Cys-2 to Asn-10.
876012	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5590 as residues: Trp-30 to Thr-43.
876018	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5592 as residues: Pro-52 to Asn-63, Pro-70 to Ile-79, Arg-93 to Gln-111.
876021	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5594 as residues: Ala-59 to Ser-72, Ser-84 to Leu-94, Thr-98 to Lys-105, Val-109 to Glu-119, Asn-124 to Leu-139, Pro-146 to Ala-155, Ser-161 to Thr-190, Glu-216 to His-221, Asn-229 to Gly-240, Ile-258 to Gly-269, Thr-300 to Thr-310, Thr-312 to Ser-317.
876022	Preferred epitopes include those comprising a sequence shown in SEQ

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	ID NO. 5595 as residues: Leu-2 to Tyr-11, Glu-55 to Thr-60.
976022	
876023	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5596 as residues: Lys-45 to Phe-58, Pro-99 to Gly-105, Arg-124
07/00/4	to Arg-130.
876024	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5597 as residues: Cys-7 to Arg-12, Pro-32 to Ser-49, Arg-59 to
	Gly-70, Ala-74 to Arg-82.
876028	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5601 as residues: Gly-46 to Gly-51.
876029	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5602 as residues: Ala-4 to Thr-9, Gln-17 to Thr-40.
876044	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5606 as residues: Asn-6 to Lys-12, His-32 to Phe-41.
876045	Preferred epitopes include those comprising a sequence shown in SEQ
	1D NO. 5607 as residues: Thr-5 to Glu-14, Pro-23 to Tyr-28, Arg-42 to
	Pro-49, Lys-87 to Ser-95.
876048	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5608 as residues: Gln-1 to Asp-11, Arg-18 to Gly-23, Thr-31 to
	Pro-38.
876057	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5611 as residues: Glu-17 to Ser-42.
876059	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5612 as residues: Pro-34 to His-49.
876065	Preferred epitopes include those comprising a sequence shown in SEQ
1	ID NO. 5614 as residues: Ser-28 to Val-33, Gln-41 to Gln-46, Gln-53 to
	Gln-63, Ala-76 to His-84, lle-88 to Ser-93, Pro-99 to Ala-105, Pro-114
	to Ser-122, Pro-145 to Thr-153, Pro-197 to Gln-206.
876078	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5616 as residues: Arg-71 to Trp-80, Arg-88 to Arg-99.
876079	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5617 as residues: Cys-16 to His-21, Lys-23 to Asp-31.
876081	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5618 as residues: Pro-6 to Cys-12.
876086	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5620 as residues: Cys-66 to Ser-74, Arg-81 to His-90.
876089	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5622 as residues: Ser-2 to Gly-11.
876090	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5623 as residues: Gln-1 to Glu-13, Lys-25 to Ser-34, Asp-49 to
	Gln-54.
876091	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5624 as residues: Phe-14 to Tyr-19, Arg-24 to Arg-32.
876093	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5625 as residues: Ser-1 to Glu-8, Asp-30 to Gly-37, Val-44 to
	Glu-58.
876094	Preferred epitopes include those comprising a sequence shown in SEQ
","	ID NO. 5626 as residues: Gly-1 to Gly-7, Ile-23 to Ala-29, Phe-40 to
1	Gin-45.
876095	Preferred epitopes include those comprising a sequence shown in SEQ
070033	1 referred chitopes metide those comprising a sequence shown in SEQ

ID NO. 5627 as residues: Lys-1 to Lys-6, Pro-8 to Glu-19. 876097 Preferred epitopes include those comprising a sequence shown in SE ID NO. 5628 as residues: Arg-30 to Ser-37.	Q
ID NO. 5628 as residues: Arg-30 to Ser-37.	•
876098 Preferred epitopes include those comprising a sequence shown in SE	Q
ID NO. 5629 as residues: Leu-18 to Leu-23.	
876101 Preferred epitopes include those comprising a sequence shown in SE	Q
ID NO. 5630 as residues: Gly-56 to Asp-62.	
876104 Preferred epitopes include those comprising a sequence shown in SE	Q _
ID NO. 5631 as residues: Gln-1 to Glu-7, Ala-31 to Glu-48.	
876107 Preferred epitopes include those comprising a sequence shown in SE	
ID NO. 5633 as residues: Gly-13 to Gln-19, Arg-58 to Gly-63, Leu-	29
to Pro-134.	
876118 Preferred epitopes include those comprising a sequence shown in SE	
ID NO. 5637 as residues: Pro-35 to Gly-42, Pro-62 to Arg-74, Val-8	7 to
Ala-93, Leu-119 to Ala-124.	
Preferred epitopes include those comprising a sequence shown in SE	
ID NO. 5638 as residues: Pro-2 to Pro-35, Ser-40 to Leu-57, Thr-83	
Thr-93, His-96 to Thr-105, Leu-114 to Gly-125, Asp-128 to Asp-133 Lys-146 to Ser-156.	,
876140 Preferred epitopes include those comprising a sequence shown in SE	<u> </u>
ID NO. 5644 as residues: Ala-39 to Leu-47, Val-49 to Lys-55, Thr-6	
to Asp-75, Thr-85 to Gly-104, Ala-114 to Gly-147, Pro-176 to Thr-1	
Ser-205 to Ser-221, Glu-233 to Lys-240, Lys-246 to Asp-251, Glu-23	
to Ser-267, Ser-291 to Leu-302, Thr-305 to Asp-324, Cys-336 to Val	
345, Phe-367 to Cys-375.	
876151 Preferred epitopes include those comprising a sequence shown in SE	Q
ID NO. 5648 as residues: Gly-101 to Arg-106.	
876152 Preferred epitopes include those comprising a sequence shown in SE	Q
ID NO. 5649 as residues: Arg-1 to Gly-12, His-33 to Leu-42.	
876155 Preferred epitopes include those comprising a sequence shown in SE	Q
ID NO. 5651 as residues: Phe-26 to Lys-51, Gln-61 to Asp-75, Gly-	
to Asn-92, Asn-101 to Cys-106, Lys-119 to Leu-124, Pro-126 to Tyr	
135, Ser-137 to Ser-150, His-161 to Ser-168, Asp-175 to Ser-182, As	n-
189 to Lys-207, Pro-225 to Thr-234, His-240 to Gly-259, Glu-266 to Val-271, Asp-285 to Ala-290, Asn-321 to Ile-353.	
876156 Preferred epitopes include those comprising a sequence shown in SE	$\overline{}$
ID NO. 5652 as residues: Lys-21 to Gly-26.	Y
876170 Preferred epitopes include those comprising a sequence shown in SE	<u> </u>
ID NO. 5656 as residues: Arg-15 to Arg-21.	Y
876172 Preferred epitopes include those comprising a sequence shown in SE	$\overline{}$
ID NO. 5657 as residues: Trp-73 to Trp-80, Tyr-90 to Lys-97, Lys-1	
to Trp-111.	
876174 Preferred epitopes include those comprising a sequence shown in SE	$\overline{\mathbf{z}}$
ID NO. 5658 as residues: Gly-7 to Glu-12, Ser-16 to Gln-25.	`
876177 Preferred epitopes include those comprising a sequence shown in SE	$\overline{\mathfrak{d}}$
ID NO. 5659 as residues: Phe-9 to Tyr-15.	[
876182 Preferred epitopes include those comprising a sequence shown in SE	
ID NO. 5661 as residues: Pro-28 to Arg-34, His-66 to Pro-81, Ser-83	to
Ala-93, Gly-98 to Lys-114.	

876184	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5663 as residues: Asn-35 to Cys-40, Ser-75 to Phe-84.
876192	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5665 as residues: Thr-4 to Ser-14, Ile-83 to Ala-94.
876198	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5667 as residues: Pro-7 to Thr-17.
876200	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5668 as residues: Leu-43 to Pro-50.
876201	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5669 as residues: Pro-28 to Glu-37.
876206	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5670 as residues: Gly-29 to Asp-39.
876207	Preferred epitopes include those comprising a sequence shown in SEQ
•	ID NO. 5671 as residues: Arg-54 to Lys-95.
876208	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5672 as residues: Ser-44 to Leu-49, Lys-52 to Pro-57, Gly-65 to
	Phe-71, Asp-94 to Trp-99, Gly-137 to Asp-149, Ser-154 to Glu-159,
	Glu-168 to Ile-173.
876209	Preferred epitopes include those comprising a sequence shown in SEQ
07.501.5	ID NO. 5673 as residues: Gly-101 to Arg-107, Ser-112 to Cys-117.
876215	Preferred epitopes include those comprising a sequence shown in SEQ
07/02/	ID NO. 5675 as residues: Phe-27 to Ile-34.
876224	Preferred epitopes include those comprising a sequence shown in SEQ
İ	ID NO. 5677 as residues: Ser-58 to Gly-63, Thr-69 to Gly-76, Ser-107 to Thr-115, Ser-144 to Gly-151, Leu-175 to Trp-181.
876226	Preferred epitopes include those comprising a sequence shown in SEQ
870220	ID NO. 5678 as residues: Arg-57 to Thr-62.
876228	Preferred epitopes include those comprising a sequence shown in SEQ
0.0220	ID NO. 5679 as residues: Glu-7 to Ser-25, Lys-39 to Leu-46.
876229	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5680 as residues: Phe-48 to Ser-58.
876232	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5681 as residues: Thr-3 to Thr-8.
876238	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5683 as residues: Asn-30 to Lys-43, Pro-58 to Glu-65, Arg-77
	to Asn-85.
876239	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5684 as residues: Thr-7 to Pro-15.
876259	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5685 as residues: Lys-1 to Gln-7, Gly-39 to Ile-50, Ile-68 to
	Cys-84, Leu-92 to Glu-99, Glu-109 to Glu-121, Pro-156 to Cys-172,
07/2/2	Pro-174 to Thr-189, Arg-212 to Gln-227.
876260	Preferred epitopes include those comprising a sequence shown in SEQ
976261	ID NO. 5686 as residues: Ala-40 to Ala-45.
876261	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5687 as residues: Arg-18 to Thr-31, Ala-39 to Gly-50, Ser-71 to Val-76.
876265	Preferred epitopes include those comprising a sequence shown in SEQ
0/0203	ID NO. 5688 as residues: Thr-4 to Ser-9.
L	1D 110. 5000 as residues. 111-4 to 501-5.

876266	Preferred epitopes include those comprising a sequence shown in SEQ
27 (272	ID NO. 5689 as residues: Leu-26 to Lys-39.
876270	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5691 as residues: Pro-20 to Arg-27.
876274	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5692 as residues: Asn-52 to Ile-58.
976277	
876277	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5694 as residues: Arg-21 to Arg-30.
876280	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5696 as residues: His-16 to Phe-21.
876281	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5697 as residues: Gln-1 to Ser-8, Val-41 to Arg-47.
876282	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5698 as residues: Gln-1 to Val-6, Asp-8 to Thr-16.
876284	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5699 as residues: Ala-24 to Arg-30, Thr-88 to Pro-107.
876306	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5702 as residues: Gly-1 to Val-9, Pro-47 to His-57.
876308	
8/0308	Preferred epitopes include those comprising a sequence shown in SEQ
1	ID NO. 5703 as residues: Lys-28 to Ser-42, Gln-49 to Lys-57, Ser-76 to
	Gly-83, Glu-99 to Val-106, Gln-132 to His-142, Ala-202 to Trp-210,
	His-271 to Ile-287.
876309	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5704 as residues: His-58 to Ala-63, Arg-86 to Gly-92.
876322	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5705 as residues: Pro-33 to Arg-38, Thr-82 to Asp-88, Ala-103
	to Lys-111, Lys-117 to Phe-122.
876326	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5706 as residues: Ser-15 to Asp-28, Glu-37 to Gly-42.
876330	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5708 as residues: Arg-41 to Lys-56.
876335	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5711 as residues: Glu-8 to Cys-16, Pro-22 to Gln-32, Lys-40 to
	Pro-49.
876340	Preferred epitopes include those comprising a sequence shown in SEQ
0,0540	ID NO. 5712 as residues: Pro-1 to Glu-18, Gly-26 to Pro-33, Pro-66 to
	Gly-75.
876345	Preferred epitopes include those comprising a sequence shown in SEQ
670343	· · · · · · · · · · · · · · · · · · ·
876354	ID NO. 5713 as residues: Arg-1 to Gly-10.
876354	Preferred epitopes include those comprising a sequence shown in SEQ
075051	ID NO. 5714 as residues: Pro-12 to Thr-18.
876361	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5715 as residues: Arg-14 to Val-29.
876364	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5716 as residues: Gln-22 to Gly-28.
876370	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5717 as residues: Gly-4 to Arg-12, Gly-33 to Cys-46.
876372	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5718 as residues: Lys-30 to Glu-35.

876374	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5719 as residues: Ser-2 to Ser-8, Glu-26 to His-33, Ser-56 to Gly-61.
876380	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5722 as residues: Ser-11 to Pro-16.
876382	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5724 as residues: Glu-15 to Ser-20.
876383	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5725 as residues: Tyr-16 to Thr-21, Lys-33 to Gln-39.
876385	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5726 as residues: Leu-11 to Phe-16.
876395	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5729 as residues: Arg-7 to Ser-26.
876397	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5730 as residues: Pro-19 to Gln-25, Thr-41 to Pro-47.
876398	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5731 as residues: Glu-1 to Arg-7.
876400	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5733 as residues: Gln-13 to Trp-20, Gly-60 to Phe-65, Cys-69 to Trp-77.
876401	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5734 as residues: Gly-25 to Trp-30, Arg-37 to Gly-44, Ser-46 to Arg-59, Ser-70 to Ser-76, Leu-106 to Gly-112.
876404	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5736 as residues: Tyr-1 to Gly-17.
876405	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5737 as residues: Tyr-1 to Ala-6, Trp-30 to Ser-36, Asp-48 to Ile-62, Ile-91 to Ile-100, Asn-119 to Asn-128, Glu-146 to Glu-152.
876408	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5738 as residues: Gly-7 to Leu-15.
876409	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5739 as residues: Gly-10 to Asn-15.
876418	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5740 as residues: Pro-57 to Asp-63.
876420	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5742 as residues: Pro-6 to Ser-12.
876426	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5745 as residues: Phe-2 to Thr-12.
876428	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5747 as residues: Thr-4 to Trp-10, Pro-25 to Ala-31.
876431	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5748 as residues: Thr-1 to Gln-6, Lys-15 to Glu-23, Pro-39 to Ile-44, Asn-63 to Gln-71, Gln-101 to Arg-111, Leu-118 to Ser-124, Leu-141 to Val-146, Pro-154 to Pro-161, Ser-187 to Pro-192, Arg-207 to Met-245, Ala-253 to Ser-263.
876432	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5749 as residues: Lys-45 to Asn-55.
876435	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5750 as residues: Asp-84 to Asn-91.

876436	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5751 as residues: Pro-81 to His-89.
876440	Preferred epitopes include those comprising a sequence shown in SEQ
075115	ID NO. 5752 as residues: Asp-1 to Leu-6, Glu-55 to Ser-60.
876441	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5753 as residues: Pro-14 to Leu-21, Cys-34 to Gly-39.
876448	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5756 as residues: Thr-1 to Glu-11, Thr-19 to Lys-30, Asn-32 to
	Glu-39, Leu-60 to Tyr-111, Ala-127 to Phe-132, Pro-184 to Thr-306.
876451	Description of the state of the
870431	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5757 as residues: Thr-52 to Lys-59.
876452	Preferred epitopes include those comprising a sequence shown in SEQ
0,0132	ID NO. 5758 as residues: Asn-1 to Arg-11, Val-23 to Ser-28, Asp-35 to
	The 40 Gly 116 to Arg 122 Levy 162 to Ser 170 Us 267 to G
876464	Thr-40, Glu-116 to Arg-122, Leu-163 to Ser-170, Ile-267 to Ser-272.
870404	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5761 as residues: Thr-6 to Lys-11, Pro-58 to Ile-72, Ser-81 to
876465	Gly-94.
876465	Preferred epitopes include those comprising a sequence shown in SEQ
076460	ID NO. 5762 as residues: Pro-2 to Trp-11, Pro-26 to Ala-32.
876469.	Preferred epitopes include those comprising a sequence shown in SEQ
25 (15)	ID NO. 5763 as residues: Trp-1 to Leu-17.
876470	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5764 as residues: Pro-30 to Glu-41, Cys-62 to Trp-68, Leu-78
	to Asn-97, Arg-131 to Asn-136.
876471	Preferred epitopes include those comprising a sequence shown in SEQ
<u></u>	ID NO. 5765 as residues: Val-7 to Leu-13, Glu-26 to Gln-32.
876472	Preferred epitopes include those comprising a sequence shown in SEQ
256155	ID NO. 5766 as residues: Ser-91 to Gly-101.
876473	Preferred epitopes include those comprising a sequence shown in SEQ
076476	ID NO. 5767 as residues: His-12 to His-22.
876476	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5770 as residues: Phe-2 to Trp-7, Cys-35 to Asn-46, Pro-55 to
	Asn-70, Pro-131 to Cys-137, Phe-141 to Thr-154, Ala-166 to Phe-177.
876481	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5772 as residues: Ala-87 to Ser-94, Asp-104 to Arg-112, Leu-
	114 to Asp-119, Ser-186 to Thr-202.
876483	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5773 as residues: Gly-1 to Pro-6.
876484	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5774 as residues: Met-2 to Leu-9, Lys-11 to Pro-28, Asp-57 to
	Leu-68, Gln-81 to Phe-118.
876487	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5775 as residues: Lys-1 to Ser-7.
876490	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5776 as residues: Glu-12 to Asp-17, Thr-26 to His-34, Asn-48
	to Tyr-57.
876491	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5777 as residues: Arg-1 to Gln-11.
876494	Preferred epitopes include those comprising a sequence shown in SEQ
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	ID NO. 5778 as residues: Asn-40 to Thr-45, His-210 to Pro-215, Glu-
	369 to Thr-375, Lys-383 to Leu-397, Pro-438 to Ile-447, Pro-510 to Tyr-
	520, Arg-528 to Arg-533, Thr-549 to Thr-555.
876495	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5779 as residues: Arg-11 to Arg-29, Arg-99 to Gly-105.
876496	Preferred epitopes include those comprising a sequence shown in SEQ
0,0.50	ID NO. 5780 as residues: Glu-1 to Gly-10.
876498	Preferred epitopes include those comprising a sequence shown in SEQ
0,0150	ID NO. 5781 as residues: Ser-1 to Ser-14.
876499	Preferred epitopes include those comprising a sequence shown in SEQ
070477	ID NO. 5782 as residues: Pro-19 to Tyr-25.
876504	Preferred epitopes include those comprising a sequence shown in SEQ
870304.	ID NO. 5784 as residues: His-7 to Asp-12, Glu-21 to Lys-26, Gln-79 to
	Ser-87.
876507	Preferred epitopes include those comprising a sequence shown in SEQ
870307	ID NO. 5785 as residues: Pro-1 to Ser-12, Leu-26 to Gly-54, Thr-61 to
	Ala-73.
876513	Preferred epitopes include those comprising a sequence shown in SEQ
870313	ID NO. 5787 as residues: Ser-3 to Gly-39, Trp-89 to Asp-96, Glu-103
	to Asn-111, Leu-138 to Pro-145.
876518	Preferred epitopes include those comprising a sequence shown in SEQ
870318	ID NO. 5788 as residues: Met-31 to Pro-38.
876524	Preferred epitopes include those comprising a sequence shown in SEQ
870324	ID NO. 5789 as residues: Pro-26 to Gln-32.
876526	Preferred epitopes include those comprising a sequence shown in SEQ
870320	ID NO. 5790 as residues: Met-7 to Tyr-13.
876530	Preferred epitopes include those comprising a sequence shown in SEQ
8/0330	ID NO. 5791 as residues: Tyr-37 to Val-45.
876533	Preferred epitopes include those comprising a sequence shown in SEQ
6/0333	ID NO. 5792 as residues: Lys-41 to Lys-47, His-52 to Gln-58, Gln-100
	to Cys-106.
876535	Preferred epitopes include those comprising a sequence shown in SEQ
8/0333	ID NO. 5794 as residues: Asp-1 to Asp-12.
876536	Preferred epitopes include those comprising a sequence shown in SEQ
870330	ID NO. 5795 as residues: Gly-11 to Gly-28, Glu-35 to Ala-40, Leu-42
ļ	to Gly-51, Ser-65 to Cys-70.
876538	Preferred epitopes include those comprising a sequence shown in SEQ
870336	ID NO. 5796 as residues: Tyr-5 to Thr-12.
876543	Preferred epitopes include those comprising a sequence shown in SEQ
0/0343	ID NO. 5798 as residues: Gln-1 to Ala-9, Cys-56 to Gly-61, Trp-105 to
	Thr-110, Arg-150 to Thr-155, Leu-189 to Lys-195.
876544	Preferred epitopes include those comprising a sequence shown in SEQ
0/0344	ID NO. 5799 as residues: Thr-15 to Asp-27.
076545	Preferred epitopes include those comprising a sequence shown in SEQ
876545	ID NO. 5800 as residues: Arg-1 to Asp-7, Leu-19 to Lys-33, Ser-45 to
	Thr-54.
976546	
876546	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5801 as residues: Thr-15 to Lys-25, Pro-35 to Phe-42, Glu-58 to
	Thr-72, Glu-115 to Met-126, Gln-131 to Thr-139, Ser-142 to Glu-157,
	1111-72, Glu-113 to Met-120, Gli-131 to 1111-139, Set-142 to Glu-137,

	Pro-165 to Gln-188, Phe-284 to Lys-301.
876553	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5805 as residues: Arg-14 to Arg-19, Asn-27 to Val-32, Glu-68
	to Thr-77, Gly-85 to Asp-90, Asp-221 to Gln-229, Thr-236 to Val-242,
	Thr-259 to Trp-266, Ser-268 to Asn-273, Asn-283 to Gly-288.
876558	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5807 as residues: Arg-22 to Gln-34.
876559	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5808 as residues: Asn-15 to Ser-20, Arg-100 to Phe-107, Glu-
	111 to Asp-118, Ile-122 to Val-127, Cys-219 to Val-227.
876560	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5809 as residues: Pro-7 to Ser-14, Thr-26 to Cys-51, Leu-55 to
	Tyr-64.
876572	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5810 as residues: Lys-16 to Lys-21.
876575	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5811 as residues: Pro-10 to Trp-19, Glu-47 to Gly-52, Tyr-75 to
	Gly-88, Met-119 to Asp-131.
876579	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5813 as residues: Ser-2 to Pro-21.
876581	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5815 as residues: Gly-32 to Gly-44, Pro-52 to Cys-60, Asp-63
	to Leu-68, Lys-148 to Asn-160.
876583	Preferred epitopes include those comprising a sequence shown in SEQ
27.500	ID NO. 5816 as residues: Glu-19 to Cys-30.
876595	Preferred epitopes include those comprising a sequence shown in SEQ
25 (50 (ID NO. 5821 as residues: Asn-1 to Arg-8, Glu-64 to Thr-70.
876596	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5822 as residues: Lys-61 to His-66, Glu-70 to Tyr-78, Pro-90 to
	Ile-95, Val-118 to Asp-127, Asp-192 to Phe-199, Asn-274 to Met-279,
	Ser-281 to Arg-291, Thr-306 to Tyr-315, Lys-338 to Gln-343, Lys-350
	to Asp-356, Pro-374 to Asp-380, Gly-398 to Pro-405, Lys-438 to Asn-446.
876597	
870397	Preferred epitopes include those comprising a sequence shown in SEQ
876600	ID NO. 5823 as residues: His-1 to Ser-6, Glu-14 to Gly-22.
870000	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5824 as residues: Asp-22 to Pro-30, Ser-49 to Asn-57, Thr-76 to Ala-91.
876601	
070001	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5825 as residues: Leu-31 to Ser-41.
876602	
0,0002	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5826 as residues: Leu-11 to Arg-19, Arg-33 to Ala-38, Ala-40
	to Gln-46, Pro-57 to Gly-62, Ser-70 to Arg-76, Thr-97 to Arg-103, Lys-119 to Lys-124.
876608	Preferred epitopes include those comprising a sequence shown in SEQ
370008	ID NO. 5827 as residues: Val-10 to Gln-18.
876609	Preferred epitopes include those comprising a sequence shown in SEQ
670003	ID NO. 5828 as residues: Leu-39 to Gln-52.
876610	
070010	Preferred epitopes include those comprising a sequence shown in SEQ

	ID NO. 5829 as residues: Ser-11 to Glu-20.
876612	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5830 as residues: Lys-1 to Asn-8, Glu-10 to Thr-15, Ser-22 to
	Gly-28, Pro-49 to His-54.
876622	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5832 as residues: Pro-46 to Leu-51.
876630	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5833 as residues: Gln-41 to Pro-46.
876633	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5835 as residues: Ala-1 to Leu-9, Ala-48 to Asp-55.
876638	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5837 as residues: Gln-1 to Arg-12, Asp-22 to Pro-44, Lys-52 to
	Asp-62, Pro-68 to Lys-93, Pro-99 to Pro-129, Ala-138 to Ser-150, Lys-
	156 to Val-194, Ile-197 to Glu-210, Ala-213 to Ala-287, Leu-289 to
	Lys-327, Lys-330 to Gly-340, Asp-344 to Gln-360, Ile-396 to Thr-401,
	Lys-409 to Asp-418, Met-450 to Ala-460, Glu-468 to Asp-481, Ala-490
	to Ser-517, Asp-523 to Ser-555.
876643	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5838 as residues: Gln-1 to Ser-13.
876645	Preferred epitopes include those comprising a sequence shown in SEQ
1	ID NO. 5839 as residues: Gly-1 to Gln-20, Gly-22 to Glu-27, Arg-46 to
	Phe-52, Thr-64 to His-72, Pro-94 to Lys-109, Ser-143 to Ser-151.
876646	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5840 as residues: Ser-29 to Glu-34.
876647	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5841 as residues: Trp-41 to Ser-46, Glu-59 to Lys-66, Lys-75 to
	His-80.
876652	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5844 as residues: Phe-23 to Val-42.
876656	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5845 as residues: Ser-38 to Cys-51, Asn-93 to Asp-100.
876657	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5846 as residues: Pro-112 to Gly-118.
876660	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5847 as residues: Glu-20 to Arg-26, Leu-30 to Cys-36, Gln-49
1	to Ser-55, Lys-82 to Thr-90.
876666	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5848 as residues: Val-39 to Asn-46, Ser-95 to Asp-101, Lys-
	118 to Val-124.
876677	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5851 as residues: Asn-1 to Val-6, Phe-76 to Tyr-83, Gly-129 to
1	Gln-135, Thr-145 to Asp-153, Ser-210 to Gln-220, Thr-230 to Asn-236,
	Lys-242 to Ala-248.
876680	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5852 as residues: Ser-1 to Thr-9, Ala-32 to Asn-37, Thr-40 to
	Tyr-49, Gln-71 to Thr-80.
876683	Preserred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5853 as residues: Pro-18 to Gly-29, Lys-67 to Lys-89.
876685	Preferred epitopes include those comprising a sequence shown in SEQ

	ID NO. 5854 as residues: Lys-19 to Asn-25, Leu-27 to Leu-38, Val-61
	to Val-68, Leu-152 to Tyr-159, Glu-222 to Cys-228, Asp-260 to Leu-
	265
876687	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5855 as residues: Ala-60 to Arg-65, Ala-82 to Arg-87.
876689	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5856 as residues: Arg-1 to Asn-9, Gln-20 to Asn-27, His-29 to
	Arg-34.
876690	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5857 as residues: Pro-15 to Asn-25, Glu-48 to Phe-59, Ser-69 to
	Arg-74, Ala-77 to Ser-82, Leu-99 to Asn-105, Ala-108 to Pro-124, Ser-
	137 to Phe-150, Ser-173 to Gly-178, Pro-186 to Pro-191, Ala-199 to
	Lys-213, Val-229 to Asp-238, Arg-272 to Asn-290.
876693	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5858 as residues: Glu-3 to Gly-12, Arg-20 to Gln-30, Leu-34 to
	Gln-39, Asp-51 to Arg-58, Gln-69 to Val-77, Gly-105 to Lys-117.
876696	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5859 as residues: Arg-1 to Arg-7, Gly-72 to Asp-78, Lys-83 to
	Gln-90.
876701	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5861 as residues: Thr-22 to Lys-31.
876716	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5862 as residues: Tyr-28 to Leu-33, Ala-70 to Lys-87, Glu-106
	to Gly-124, Gly-127 to Glu-160, Leu-179 to Asp-194.
876719	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5863 as residues: Asn-19 to Ser-25, Gln-57 to Leu-66, Asp-76
	to Ser-81, Glu-101 to Gln-106, Phe-121 to Asp-127, Ser-133 to Asp-
076706	146, Thr-186 to Lys-197, Arg-259 to Leu-266, Asn-268 to Leu-274.
876725	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5865 as residues: Thr-23 to Pro-34, Glu-39 to Asp-83, Asn-89
	to Lys-99, Asp-118 to Asp-128, Asp-135 to Glu-150, Glu-153 to Gly-
	168, Gly-181 to Thr-187, Arg-200 to Asp-205, Arg-273 to Ile-279, Thr-295 to Asp-300, Thr-316 to Cys-321.
876726	Preferred epitopes include those comprising a sequence shown in SEQ
070720	ID NO. 5866 as residues: Tyr-17 to Gly-22, Lys-29 to Tyr-34, Asp-39
	to Asp-44, Leu-71 to Glu-76, Pro-164 to Gly-171.
876732	Preferred epitopes include those comprising a sequence shown in SEQ
0.0.52	ID NO. 5869 as residues: Ser-1 to Gln-6, Leu-57 to Phe-62, Arg-86 to
	Glu-91.
876744	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5871 as residues: Thr-98 to Ser-104, Thr-115 to Tyr-126, Gln-
	149 to Glu-164.
876745	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5872 as residues: His-1 to Gln-7, Trp-14 to Gln-29, Arg-41 to
	Pro-48, Leu-91 to His-97, Pro-99 to Ser-114, Ser-119 to Gly-124.
876747	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5873 as residues: Ala-13 to Arg-35, Pro-58 to Met-75, Asn-104
	to Ser-119, Pro-144 to Ile-167, Lys-183 to Phe-224, Cys-246 to Gly-252,
	Lys-304 to Gly-320.
876750	Preferred epitopes include those comprising a sequence shown in SEQ

	ID NO. 5874 as residues: Ala-1 to Ser-6, Ser-29 to Ser-37, Gln-52 to Tyr-58.
876752	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5875 as residues: Pro-44 to Gly-51.
876753	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5876 as residues: Arg-5 to Arg-12.
876760	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5877 as residues: Thr-11 to Ala-16, Thr-85 to Glu-92, Asn-114
	to Glu-122, Asp-150 to Gly-156.
876762	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5878 as residues: Pro-14 to Ile-24, Thr-35 to Pro-46.
876771	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5881 as residues: His-28 to Gly-33.
876773	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5882 as residues: Gly-3 to Thr-9, Glu-39 to Lys-48, Arg-134 to Lys-139, Pro-147 to Val-152, Thr-167 to Glu-172, His-190 to Gln-196.
876791	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5885 as residues: Pro-1 to Glu-20, Leu-79 to Ser-87, Lys-90 to Gly-96, Gln-109 to Thr-121, Val-133 to Gly-139.
876798	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5887 as residues: Thr-25 to Val-31, Lys-47 to Asp-62.
876802	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5888 as residues: Leu-2 to Thr-8, Asp-15 to Gly-26, Phe-64 to Ser-70, Pro-77 to Trp-82, Pro-85 to Lys-90.
876807	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5890 as residues: Lys-12 to Ser-18, Tyr-26 to Thr-33, Leu-71 to Thr-76, Pro-102 to Ser-110, Asp-114 to Gln-119, Glu-137 to Asp-159, Gly-162 to His-172, Thr-179 to Gly-194, Ala-198 to Asp-229.
876809	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5891 as residues: Arg-7 to Lys-13, Pro-28 to Cys-34, Gly-100 to Asn-109, Cys-155 to Arg-162, Glu-214 to Gln-219, Glu-240 to Asp-246, Gly-254 to His-265.
876817	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5894 as residues: Pro-22 to Asn-28, Pro-47 to Asn-57, Glu-92 to Gly-98, Pro-120 to Ile-135, Ala-138 to Cys-155, Pro-161 to Val-181, Ala-185 to Asp-196, Val-207 to Asn-213, Asn-219 to Asn-236, Asn-242 to Asn-250, Leu-252 to Asn-274, Ala-281 to Cys-295, Pro-297 to Cys-311, Pro-317 to Asn-339, Thr-417 to Tyr-423, Gln-443 to Gly-458, Thr-471 to His-476, Thr-484 to Gln-490, Asp-497 to Trp-511.
876823	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5896 as residues: Arg-1 to Trp-23, Pro-37 to Gly-47, Gly-50 to His-56, Phe-64 to Gly-74, Pro-76 to Ala-81, Pro-84 to Gly-95, Pro-101 to Pro-112, Lys-135 to Lys-146, Lys-159 to Asp-176.
876829	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5897 as residues: Pro-51 to His-56, Glu-69 to Asn-74, Gly-190 to Lys-199.
876830	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5898 as residues: Asp-27 to Gly-39.
876842	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5902 as residues: Glu-8 to Arg-13, Leu-17 to Val-23.

876856	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5903 as residues: Glu-63 to Asn-73, Pro-114 to Tyr-122, Ser-194 to Glu-201, Ile-263 to Ser-269.
876858	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5904 as residues: Asn-1 to Val-6, Lys-9 to Gln-16, Asn-47 to Glu-53, Asn-116 to Ser-121, Pro-130 to Thr-139, His-159 to Glu-165.
876865	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5905 as residues: Leu-26 to Asp-39, Asp-47 to Arg-54, Glu-62 to Val-72.
876866	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5906 as residues: Ser-1 to Gln-8, Val-40 to Ser-49, Arg-105 to Lys-110.
876870	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5907 as residues: Ser-25 to Trp-32.
876873	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5908 as residues: Gln-21 to Met-26, Gln-50 to Lys-61.
876876	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5909 as residues: Ala-8 to Arg-14, Ile-64 to Thr-69, Val-94 to Asp-101, His-112 to Gln-117, Tyr-139 to Glu-145, Tyr-195 to Cys-208, Gly-216 to Gly-223, Asp-297 to Ser-307, Gly-378 to Leu-383, Ile-391 to Pro-404, Asn-451 to Ser-466.
876878	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5910 as residues: Pro-32 to Arg-41.
876882	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5911 as residues: Thr-4 to Gly-13, Asp-20 to Val-25, Ala-46 to Asn-65, Gly-69 to Gly-75, Pro-82 to Gly-113, Pro-119 to Pro-174, Gly-181 to Gly-187, Tyr-190 to Tyr-212.
876886	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5912 as residues: Ser-9 to Arg-22, Gln-28 to Trp-34, Gly-36 to Leu-43, Arg-45 to Trp-52, Asp-56 to Leu-61, Ala-65 to Tyr-72, Leu-102 to Gly-109, Pro-111 to Ala-116, Ala-120 to Arg-125, His-129 to Gln-134, Pro-136 to Gly-145, Pro-167 to Thr-172, Glu-232 to Lys-239, Lys-253 to Asn-258, Leu-357 to Gly-362, Leu-371 to Gly-376.
876888	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5913 as residues: Glu-31 to Asp-39.
876890	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5914 as residues: Glu-91 to Pro-100, Tyr-122 to Thr-127, Thr-168 to Val-173, Thr-210 to Asp-215, Leu-219 to Gly-224, Gly-232 to Val-237.
876892	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5915 as residues: Ser-8 to Ser-20.
876901	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5916 as residues: Tyr-130 to Glu-136, Arg-148 to His-159, Pro-214 to Leu-221, His-224 to Gly-229, Glu-238 to Glu-246, Gln-331 to Trp-343.
876904	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5918 as residues: Val-61 to Gln-69, Gln-106 to Thr-111.
876905	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5919 as residues: Arg-1 to Arg-7, Pro-29 to Lys-56, Asp-103 to Arg-108, Tyr-122 to Ser-127, Gly-219 to Glu-227, Asp-250 to Glu-255,

	Glu-294 to Pro-301, Ala-321 to Tyr-327, Arg-367 to Pro-373, Glu-396 to Asn-405, Gly-411 to Arg-418, Asn-433 to Lys-441.
876909	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5920 as residues: Ala-32 to Ala-40, Glu-93 to Phe-103, Lys-173 to Thr-189.
876912	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5921 as residues: Glu-40 to Pro-47, Lys-56 to Trp-62.
876920	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5923 as residues: Arg-1 to Gly-15, Ser-42 to Trp-51, Pro-59 to Arg-64.
876921	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5924 as residues: Tyr-1 to Leu-6.
876923	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5925 as residues: Pro-6 to Cys-14, Glu-33 to Leu-38, Val-209 to Lys-216, Pro-270 to Gln-278, His-321 to Thr-330.
876936	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5928 as residues: Ala-54 to His-67, Pro-69 to Lys-86.
876940	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5930 as residues: Ala-1 to Asp-29, Pro-51 to His-59, Asn-67 to Asp-73.
876941	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5931 as residues: Pro-16 to Arg-28.
876942	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5932 as residues: Glu-1 to Gln-6, Val-8 to Trp-15.
876943	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5933 as residues: Gly-1 to Gln-9, Asn-11 to Arg-16, Cys-28 to His-33, Pro-51 to Pro-57, Glu-66 to Glu-72, Pro-84 to Asp-89, Pro-104 to Asp-109, Glu-122 to Thr-132.
876944	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5934 as residues: Arg-3 to Gly-11.
876945	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5935 as residues: Pro-15 to Pro-24.
876946	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5936 as residues: Ser-8 to Ser-14.
876947	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5937 as residues: Gly-27 to Ala-34.
876949	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5938 as residues: Pro-5 to His-14, Arg-38 to Gln-43, Leu-80 to Arg-86.
876952	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5939 as residues: Ser-8 to Thr-18, Pro-52 to Ala-61, Pro-67 to Gly-72, Ala-81 to Thr-88, Glu-105 to Thr-120.
876953	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5940 as residues: Gly-1 to Asp-12, Ser-64 to Trp-74, Met-82 to Tyr-88, Phe-101 to Cys-106, Tyr-120 to Lys-132.
876954	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5941 as residues: Pro-1 to Ile-12, Asp-30 to Tyr-35, Leu-38 to Pro-45, Lys-54 to Thr-60, Thr-75 to Leu-80, Asp-92 to Tyr-100, Ile-133 to Thr-138, Thr-194 to Glu-199, Asp-233 to Leu-239, Met-243 to Ala-

	251, Asp-254 to Glu-261.
876957	Preferred epitopes include those comprising a sequence shown in SEQ
870557	ID NO. 5942 as residues: Lys-71 to Asn-88, Ala-115 to Cys-130, Ala-
	175 to Arg-182.
876958	
070938	Preferred epitopes include those comprising a sequence shown in SEQ
976062	ID NO. 5943 as residues: Gln-1 to Pro-8.
876963	Preferred epitopes include those comprising a sequence shown in SEQ
976064	ID NO. 5946 as residues: Val-16 to Ser-21, Ala-60 to Lys-72.
876964	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5947 as residues: Thr-6 to Lys-13, Met-16 to Glu-36, Lys-59 to
076066	Phe-65, Leu-71 to Gln-77.
876966	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5949 as residues: Lys-13 to Trp-19, Ser-25 to Gln-32, Glu-53 to
25.00	Gln-58, Arg-108 to Gly-113.
876967	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5950 as residues: Lys-1 to Asp-9, Arg-16 to Gly-21, Cys-51 to
	Val-59, Asp-65 to Ser-71, Thr-79 to Asn-90, Asn-99 to Asn-111, Ser-
	149 to Pro-156.
876968	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5951 as residues: Asn-44 to Tyr-49, Gly-71 to Glu-79.
876969	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5952 as residues: Arg-74 to Arg-79.
876975	Preferred epitopes include those comprising a sequence shown in SEO
	ID NO. 5954 as residues: Phe-12 to Ile-19, Arg-25 to Arg-31.
876976	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5955 as residues: Asn-78 to Gln-92.
876977	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5956 as residues: Asn-1 to Glu-8, Ala-38 to Gly-46, Gln-58 to
	Asp-71, Ala-75 to Cys-103, Met-106 to Ala-140, Gln-153 to Ile-159.
876981	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5959 as residues: Gln-40 to Lys-45.
876983	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5960 as residues: Leu-37 to Pro-42.
876984	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5961 as residues: His-5 to Thr-11, Arg-71 to Pro-77.
876985	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5962 as residues: Tyr-7 to Gly-28, Arg-38 to Asp-65, Asp-78 to
	Ser-90, Ser-92 to Ser-115, Asp-117 to Ser-132, Val-148 to Leu-153,
<u></u>	Lys-155 to His-168.
876987	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5963 as residues: Lys-30 to Thr-35, Ser-49 to Tyr-55.
876989	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5964 as residues: Gly-4 to Gly-10, Glu-17 to Gly-28, Met-35 to
	Asp-41, Glu-79 to Gln-85, Gln-102 to Gly-110.
876992	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5967 as residues: Ser-15 to Pro-21.
876993	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5968 as residues: His-44 to Gln-52, Pro-55 to Lys-72, Ser-87 to
	Ser-93, Arg-105 to Leu-111, Phe-119 to Lys-124.
L	1 001-75, 70 g-105 to Leu-111, File-119 to Lys-124.

876994	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5969 as residues: Leu-28 to Glu-33, Met-54 to Cys-60.
876998	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5971 as residues: Glu-1 to Pro-25, Gly-30 to Ala-54, Asn-65 to
	Asn-82, Leu-89 to Ser-97.
877000	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5972 as residues: Ala-1 to Asn-6, Val-8 to Tyr-20.
877002	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5974 as residues: Ser-32 to Gly-53, Thr-61 to Ser-70.
877005	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5976 as residues: Gly-12 to Gly-22.
877006	Preferred epitopes include those comprising a sequence shown in SEQ
077007	ID NO. 5977 as residues: Glu-8 to Ser-14, Thr-26 to Asn-40.
877007	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 5978 as residues: Glu-31 to Leu-38.
877008	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5979 as residues: Ser-37 to Ser-47, Gln-58 to Thr-69, Val-72 to
	Gln-77, Gly-125 to Lys-155.
877010	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5981 as residues: Gly-20 to Ser-29.
877011	Preferred epitopes include those comprising a sequence shown in SEQ
022014	ID NO. 5982 as residues: Ser-30 to Trp-36.
877014	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5985 as residues: Asp-1 to Arg-31, Lys-35 to Lys-44, Glu-55 to Leu-61, Thr-71 to Asp-76, Ile-82 to Asn-101.
877015	Preferred epitopes include those comprising a sequence shown in SEQ
877013	ID NO. 5986 as residues: Lys-1 to His-12, Ser-26 to Thr-31, His-54 to
Ì	Val-60.
877018	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5987 as residues: Gly-9 to Glu-16, Asn-46 to Glu-54.
877019	Preferred epitopes include those comprising a sequence shown in SEQ
1.	ID NO. 5988 as residues: Lys-24 to Glu-38, Arg-48 to Ala-54, Gly-61
	to Ala-67.
877022	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5990 as residues: Arg-10 to Gly-15, Thr-55 to Lys-64.
877024	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 5992 as residues: Thr-19 to Pro-26.
877025	Preferred epitopes include those comprising a sequence shown in SEQ
077036	ID NO. 5993 as residues: Gly-19 to Asn-27. Preferred epitopes include those comprising a sequence shown in SEQ
877026	ID NO. 5994 as residues: Met-27 to Asn-34, Val-57 to Glu-84, Glu-86
	to Ala-100, Asp-122 to Ala-128.
877027	Preferred epitopes include those comprising a sequence shown in SEQ
077027	ID NO. 5995 as residues: Gln-36 to Ser-42.
877030	Preferred epitopes include those comprising a sequence shown in SEQ
3.,030	ID NO. 5997 as residues: Glu-30 to Ala-35, Leu-39 to Ser-44, Pro-50 to
	Asp-57.
877037	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6001 as residues: Gln-61 to Lys-67.

BY/044 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6003 as residues: Apr2-22 to Gly-27, Ser-34 to Gly-39.	0.550.11	
877046 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6004 as residues: Phe-65 to Trp-73, Arg-87 to Gly-92, Gly-107 to Lys-112, Pro-177 to Trr-186, Glu-251 to Arg-256, Phe-282 to Lys-287. 877047 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6005 as residues: Try-2 to Gly-8. 877049 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6006 as residues: Pro-2 to Pro-11. 877050 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6007 as residues: Ser-36 to Lys-42. 877051 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6008 as residues: Gln-5 to Arg-12, Tyr-32 to Ser-43. 877056 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6010 as residues: Tro-52 to Val-57, Asp-59 to Gln-69. 877058 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6012 as residues: Thr-13 to Pro-20. 877059 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6013 as residues: Leu-30 to Ser-38. 877060 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6013 as residues: Leu-30 to Ser-38. 877061 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6014 as residues: Clau-30 to Ser-38. 877066 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6016 as residues: Gln-1 to Trp-11, Pro-47 to Tyr-53. 877067 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6017 as residues: Pro-32 to Asn-31, Leu-33 to Gln-29. 877070 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6020 as residues: Arg-33 to Leu-40. 877071 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6020 as residues: Pro-23 to Asn-31, Leu-33 to Phe-38. 877078 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6022 as residues: Pro-23 to Asn-31, Leu-33 to Phe-38. 877087 Preferred epitopes include those comprising a sequen	877044	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6003 as residues: Arg-22 to Gly-27, Ser-34 to Gly-39.
ID NO. 6004 as residues: Phe-65 to Trp-73, Arg-87 to Gly-92, Gly-107 to Lys-112, Pro-177 to Thr-186, Glu-251 to Arg-256, Phe-282 to Lys-287. 877047 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6005 as residues: Tyr-2 to Gly-8. 877049 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6006 as residues: Pro-2 to Pro-11. 877050 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6007 as residues: Ser-36 to Lys-42. 877051 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6008 as residues: Ser-36 to Lys-42. 877056 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6010 as residues: Pro-52 to Val-57, Asp-59 to Gln-69. 877058 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6012 as residues: Thr-13 to Pro-20. 877059 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6013 as residues: Leu-30 to Ser-38. 877063 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6014 as residues: Asp-4 to Ala-15. 877066 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6016 as residues: Gln-1 to Trp-11, Pro-47 to Tyr-53. 877067 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6016 as residues: Pro-11 to Asp-16, Arg-23 to Gln-29. 877068 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6017 as residues: Pro-11 to Asp-16, Arg-23 to Gln-29. 877070 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6018 as residues: Pro-21 to Asn-31, Leu-33 to Phe-38. 877071 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6020 as residues: Arg-33 to Leu-40. 877071 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6020 as residues: Arg-31 to Met-6, Thr-34 to Glu-54, Glu-58 to Asn-63. 877087 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6020 as residues: Pro-31 to Met-6, T	877046	
to Lys-112, Pro-177 to Thr-186, Glu-251 to Arg-256, Phe-282 to Lys-287. 877047 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6005 as residues: Tyr-2 to Gly-8. 877049 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6006 as residues: Pro-2 to Pro-11. 877050 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6007 as residues: Ser-36 to Lys-42. 877051 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6008 as residues: Gln-5 to Arg-12, Tyr-32 to Ser-43. 877056 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6010 as residues: Pro-52 to Val-57, Asp-59 to Gln-69. 877058 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6012 as residues: Thr-13 to Pro-20. 877059 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6013 as residues: Leu-30 to Ser-38. 877063 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6014 as residues: Asp-4 to Ala-15. 877066 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6016 as residues: Asp-4 to Ala-15. 877067 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6016 as residues: Gln-1 to Trp-11, Pro-47 to Tyr-53. 877067 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6016 as residues: Pro-11 to Asp-16, Arg-23 to Gln-29. 877068 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6018 as residues: Lys-26 to Arg-32. 877070 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6020 as residues: Arg-33 to Leu-40. 877071 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6020 as residues: Arg-31 to Leu-33 to Phe-38. 877073 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6022 as residues: Arg-17 to Met-6, Thr-34 to Glu-54, Glu-58 to Asn-63. 877087 Preferred epitopes include those comprising a sequence shown in SEQ ID	077010	
877047 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6005 as residues: Tyr-2 to Gly-8. 877049 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6006 as residues: Pro-2 to Pro-11. 877050 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6007 as residues: Ser-36 to Lys-42. 877051 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6008 as residues: Gln-5 to Arg-12, Tyr-32 to Ser-43. 877056 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6010 as residues: Pro-52 to Val-57, Asp-59 to Gln-69. 877058 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6012 as residues: Pro-52 to Val-57, Asp-59 to Gln-69. 877059 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6013 as residues: Leu-30 to Ser-38. 877063 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6014 as residues: Leu-30 to Ser-38. 877066 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6016 as residues: Gln-1 to Trp-11, Pro-47 to Tyr-53. 877067 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6016 as residues: Pro-11 to Asp-16, Arg-23 to Gln-29. 877068 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6018 as residues: Pro-11 to Asp-16, Arg-23 to Gln-29. 877070 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6018 as residues: Pro-11 to Asp-16, Arg-23 to Gln-29. 877071 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6020 as residues: Pro-23 to Asn-31, Leu-33 to Phe-38. 877073 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6027 as residues: Arg-33 to Leu-40. 877079 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6020 as residues: Pro-23 to Asn-31, Leu-33 to Phe-38. 877070 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6028 as residues: Thr-	1	
### BY00. 6005 as residues: Tyr-2 to Gly-8. ### B77049 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6006 as residues: Pro-2 to Pro-11. ### B77050 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6007 as residues: Ser-36 to Lys-42. ### B77051 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6008 as residues: Gln-5 to Arg-12, Tyr-32 to Ser-43. #### B77056 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6010 as residues: Pro-52 to Val-57, Asp-59 to Gln-69. #### B77058 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6012 as residues: Thr-13 to Pro-20. #### B77059 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6013 as residues: Leu-30 to Ser-38. #### B77060 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6014 as residues: Asp-4 to Ala-15. #### B77061 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6016 as residues: Gln-1 to Trp-11, Pro-47 to Tyr-53. #### B77062 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6016 as residues: Gln-1 to Trp-11, Pro-47 to Tyr-53. #### B77063 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6016 as residues: Pro-11 to Asp-16, Arg-23 to Gln-29. #### B77064 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6018 as residues: Pro-11 to Asp-16, Arg-23 to Gln-29. #### B77070 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6020 as residues: Arg-33 to Leu-40. #### B77071 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6020 as residues: Pro-23 to Asn-31, Leu-33 to Phe-38. #### B77073 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6022 as residues: Pro-23 to Asn-31, Leu-33 to Phe-38. #### B77084 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6028 as residues: Arg-17 to Gly-23. ####		
### BY00. 6005 as residues: Tyr-2 to Gly-8. ### B77049 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6006 as residues: Pro-2 to Pro-11. ### B77050 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6007 as residues: Ser-36 to Lys-42. ### B77051 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6008 as residues: Gln-5 to Arg-12, Tyr-32 to Ser-43. #### B77056 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6010 as residues: Pro-52 to Val-57, Asp-59 to Gln-69. #### B77058 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6012 as residues: Thr-13 to Pro-20. #### B77059 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6013 as residues: Leu-30 to Ser-38. #### B77060 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6014 as residues: Asp-4 to Ala-15. #### B77061 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6016 as residues: Gln-1 to Trp-11, Pro-47 to Tyr-53. #### B77062 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6016 as residues: Gln-1 to Trp-11, Pro-47 to Tyr-53. #### B77063 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6016 as residues: Pro-11 to Asp-16, Arg-23 to Gln-29. #### B77064 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6018 as residues: Pro-11 to Asp-16, Arg-23 to Gln-29. #### B77070 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6020 as residues: Arg-33 to Leu-40. #### B77071 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6020 as residues: Pro-23 to Asn-31, Leu-33 to Phe-38. #### B77073 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6022 as residues: Pro-23 to Asn-31, Leu-33 to Phe-38. #### B77084 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6028 as residues: Arg-17 to Gly-23. ####	877047	Preferred epitopes include those comprising a sequence shown in SEO
ID NO. 6006 as residues: Pro-2 to Pro-11.		ID NO. 6005 as residues: Tyr-2 to Gly-8.
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ID NO. 6007 as residues: Ser-36 to Lys-42.		ID NO. 6006 as residues: Pro-2 to Pro-11.
877051 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6008 as residues: Gln-5 to Arg-12, Tyr-32 to Ser-43. 877056 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6010 as residues: Pro-52 to Val-57, Asp-59 to Gln-69. 877058 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6012 as residues: Thr-13 to Pro-20. 877059 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6013 as residues: Leu-30 to Ser-38. 877061 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6014 as residues: Asp-4 to Ala-15. 877062 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6016 as residues: Gln-1 to Trp-11, Pro-47 to Tyr-53. 877063 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6016 as residues: Gln-1 to Trp-11, Pro-47 to Tyr-53. 877064 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6017 as residues: Pro-11 to Asp-16, Arg-23 to Gln-29. 877068 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6018 as residues: Lys-26 to Arg-32. 877070 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6020 as residues: Arg-33 to Leu-40. 877071 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6021 as residues: Pro-23 to Asn-31, Leu-33 to Phe-38. 877073 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6022 as residues: Ser-1 to Ser-17. 877087 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6027 as residues: Arg-1 to Met-6, Thr-34 to Glu-54, Glu-58 to Asn-63. 877088 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6029 as residues: Arg-1 to Gly-23. 877091 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6030 as residues: Pro-30 to Cys-43. 877094 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6030 as residues: Pro-90 to Tyr-17, Gln-29	877050	Preferred epitopes include those comprising a sequence shown in SEO
Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6008 as residues: Gln-5 to Arg-12, Tyr-32 to Ser-43. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6010 as residues: Pro-52 to Val-57, Asp-59 to Gln-69. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6012 as residues: Thr-13 to Pro-20. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6013 as residues: Leu-30 to Ser-38. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6014 as residues: Asp-4 to Ala-15. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6016 as residues: Gln-1 to Trp-11, Pro-47 to Tyr-53. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6016 as residues: Fro-11 to Asp-16, Arg-23 to Gln-29. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6018 as residues: Lys-26 to Arg-32. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6018 as residues: Arg-33 to Leu-40. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6020 as residues: Arg-33 to Leu-40. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6021 as residues: Pro-23 to Asn-31, Leu-33 to Phe-38. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6022 as residues: Ser-1 to Ser-17. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6027 as residues: Arg-1 to Met-6, Thr-34 to Glu-54, Glu-58 to Asn-63. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6028 as residues: Arg-1 to Met-6, Thr-34 to Glu-54, Glu-58 to Asn-63. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6028 as residues: Pro-33 to Cys-43. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6030 as residues: Pro-33 to Cys-43. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6031 as residues: Pro-94		
ID NO. 6008 as residues: Gln-5 to Arg-12, Tyr-32 to Ser-43. 877056 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6010 as residues: Pro-52 to Val-57, Asp-59 to Gln-69. 877058 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6012 as residues: Thr-13 to Pro-20. 877059 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6013 as residues: Leu-30 to Ser-38. 877063 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6014 as residues: Asp-4 to Ala-15. 877066 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6016 as residues: Gln-1 to Trp-11, Pro-47 to Tyr-53. 877067 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6016 as residues: Pro-11 to Asp-16, Arg-23 to Gln-29. 877068 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6018 as residues: Lys-26 to Arg-32. 877070 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6020 as residues: Arg-33 to Leu-40. 877071 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6021 as residues: Pro-23 to Asn-31, Leu-33 to Phe-38. 877073 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6021 as residues: Ser-1 to Ser-17. 877087 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6027 as residues: Arg-1 to Met-6, Thr-34 to Glu-54, Glu-58 to Asn-63. 877088 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6028 as residues: Thr-6 to Gly-13, Trp-20 to Thr-36. 877092 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6029 as residues: Pro-33 to Cys-43. 877094 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6031 as residues: Pro-33 to Cys-43. 877094 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6031 as residues: Pro-91 to Tyr-17, Gln-29 to Tyr-38, Ala-47 to Glu-55. 877096 Preferred epitopes include those comp	877051	^ ···· · · · · · · · · · · · ·
Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6010 as residues: Pro-52 to Val-57, Asp-59 to Gln-69. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6012 as residues: Thr-13 to Pro-20. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6013 as residues: Leu-30 to Ser-38. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6014 as residues: Asp-4 to Ala-15. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6016 as residues: Gln-1 to Trp-11, Pro-47 to Tyr-53. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6017 as residues: Pro-11 to Asp-16, Arg-23 to Gln-29. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6018 as residues: Lys-26 to Arg-32. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6020 as residues: Arg-33 to Leu-40. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6020 as residues: Pro-23 to Asn-31, Leu-33 to Phe-38. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6022 as residues: Pro-23 to Asn-31, Leu-33 to Phe-38. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6022 as residues: Pro-23 to Asn-31, Leu-33 to Phe-38. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6027 as residues: Arg-1 to Met-6, Thr-34 to Glu-54, Glu-58 to Asn-63. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6028 as residues: Thr-6 to Gly-13, Trp-20 to Thr-36. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6029 as residues: Pro-33 to Cys-43. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6031 as residues: Pro-9 to Tyr-17, Gln-29 to Tyr-38, Ala-47 to Glu-55. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6033 as residues: Pro-9 to Tyr-17.	077053	
### ID NO. 6010 as residues: Pro-52 to Val-57, Asp-59 to Gln-69. ### Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6012 as residues: Thr-13 to Pro-20. #### Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6013 as residues: Leu-30 to Ser-38. ##### Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6014 as residues: Asp-4 to Ala-15. ###################################	977056	
Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6012 as residues: Thr-13 to Pro-20. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6013 as residues: Leu-30 to Ser-38. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6014 as residues: Asp-4 to Ala-15. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6016 as residues: Gln-1 to Trp-11, Pro-47 to Tyr-53. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6016 as residues: Pro-11 to Asp-16, Arg-23 to Gln-29. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6018 as residues: Lys-26 to Arg-32. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6020 as residues: Arg-33 to Leu-40. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6021 as residues: Pro-23 to Asn-31, Leu-33 to Phe-38. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6022 as residues: Pro-23 to Asn-31, Leu-33 to Phe-38. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6022 as residues: Ser-1 to Ser-17. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6027 as residues: Arg-1 to Met-6, Thr-34 to Glu-54, Glu-58 to Asn-63. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6028 as residues: Arg-17 to Gly-13, Trp-20 to Thr-36. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6029 as residues: Pro-33 to Cys-43. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6031 as residues: Pro-3 to Cys-43. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6031 as residues: Pro-9 to Tyr-17, Gln-29 to Tyr-38, Ala-47 to Glu-55. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6033 as residues: Pro-9 to Tyr-17, Gln-29 to Tyr-38, Ala-47 to Glu-55.	8//056	
ID NO. 6012 as residues: Thr-13 to Pro-20.		
877059 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6013 as residues: Leu-30 to Ser-38. 877063 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6014 as residues: Asp-4 to Ala-15. 877066 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6016 as residues: Gln-1 to Trp-11, Pro-47 to Tyr-53. 877067 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6017 as residues: Pro-11 to Asp-16, Arg-23 to Gln-29. 877068 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6018 as residues: Lys-26 to Arg-32. 877070 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6020 as residues: Arg-33 to Leu-40. 877071 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6021 as residues: Pro-23 to Asn-31, Leu-33 to Phe-38. 877073 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6022 as residues: Ser-1 to Ser-17. 877087 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6022 as residues: Arg-1 to Met-6, Thr-34 to Glu-54, Glu-58 to Asn-63. 877088 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6028 as residues: Thr-6 to Gly-13, Trp-20 to Thr-36. 877092 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6029 as residues: Arg-17 to Gly-23. 877093 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6030 as residues: Pro-33 to Cys-43. 877094 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6031 as residues: Pro-9 to Tyr-17, Gln-29 to Tyr-38, Ala-47 to Glu-55. 877096 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6033 as residues: Pro-9 to Tyr-17, Gln-29 to Tyr-38, Ala-47 to Glu-55.	877058	
ID NO. 6013 as residues: Leu-30 to Ser-38.		
ID NO. 6013 as residues: Leu-30 to Ser-38.	877059	Preferred epitopes include those comprising a sequence shown in SEQ
Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6014 as residues: Asp-4 to Ala-15. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6016 as residues: Gln-1 to Trp-11, Pro-47 to Tyr-53. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6017 as residues: Pro-11 to Asp-16, Arg-23 to Gln-29. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6018 as residues: Lys-26 to Arg-32. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6020 as residues: Arg-33 to Leu-40. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6021 as residues: Pro-23 to Asn-31, Leu-33 to Phe-38. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6022 as residues: Ser-1 to Ser-17. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6027 as residues: Arg-1 to Met-6, Thr-34 to Glu-54, Glu-58 to Asn-63. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6028 as residues: Thr-6 to Gly-13, Trp-20 to Thr-36. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6029 as residues: Arg-17 to Gly-23. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6030 as residues: Pro-33 to Cys-43. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6031 as residues: Pro-3 to Cys-43. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6031 as residues: Pro-9 to Tyr-17, Gln-29 to Tyr-38, Ala-47 to Glu-55. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6031 as residues: Pro-9 to Tyr-17, Gln-29 to Tyr-38, Ala-47 to Glu-55. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6033 as residues: Pro-9 to Tyr-17, Gln-29 to Tyr-38, Ala-47 to Glu-55.		
ID NO. 6014 as residues: Asp-4 to Ala-15. 877066 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6016 as residues: Gln-1 to Trp-11, Pro-47 to Tyr-53. 877067 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6017 as residues: Pro-11 to Asp-16, Arg-23 to Gln-29. 877068 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6018 as residues: Lys-26 to Arg-32. 877070 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6020 as residues: Arg-33 to Leu-40. 877071 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6021 as residues: Pro-23 to Asn-31, Leu-33 to Phe-38. 877073 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6022 as residues: Ser-1 to Ser-17. 877087 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6027 as residues: Arg-1 to Met-6, Thr-34 to Glu-54, Glu-58 to Asn-63. 877088 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6028 as residues: Thr-6 to Gly-13, Trp-20 to Thr-36. 877092 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6029 as residues: Arg-17 to Gly-23. 877094 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6031 as residues: Pro-33 to Cys-43. 877094 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6031 as residues: Pro-9 to Tyr-17, Gln-29 to Tyr-38, Ala-47 to Glu-55. 877096 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6033 as residues: Pro-9 to Tyr-17, Gln-29 to Tyr-38, Ala-47 to Glu-55.	877063	
Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6016 as residues: Gln-1 to Trp-11, Pro-47 to Tyr-53. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6017 as residues: Pro-11 to Asp-16, Arg-23 to Gln-29. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6018 as residues: Lys-26 to Arg-32. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6020 as residues: Arg-33 to Leu-40. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6021 as residues: Pro-23 to Asn-31, Leu-33 to Phe-38. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6021 as residues: Ser-1 to Ser-17. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6027 as residues: Arg-1 to Met-6, Thr-34 to Glu-54, Glu-58 to Asn-63. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6028 as residues: Thr-6 to Gly-13, Trp-20 to Thr-36. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6029 as residues: Arg-17 to Gly-23. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6029 as residues: Pro-33 to Cys-43. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6030 as residues: Pro-33 to Cys-43. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6031 as residues: Pro-9 to Tyr-17, Gln-29 to Tyr-38, Ala-47 to Glu-55. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6031 as residues: Pro-9 to Tyr-17, Gln-29 to Tyr-38, Ala-47 to Glu-55. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6033 as residues: Pro-9 to Tyr-17, Gln-29 to Tyr-38, Ala-47 to Glu-55.		
ID NO. 6016 as residues: Gln-1 to Trp-11, Pro-47 to Tyr-53. 877067 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6017 as residues: Pro-11 to Asp-16, Arg-23 to Gln-29. 877068 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6018 as residues: Lys-26 to Arg-32. 877070 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6020 as residues: Arg-33 to Leu-40. 877071 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6021 as residues: Pro-23 to Asn-31, Leu-33 to Phe-38. 877073 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6022 as residues: Ser-1 to Ser-17. 877087 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6027 as residues: Arg-1 to Met-6, Thr-34 to Glu-54, Glu-58 to Asn-63. 877088 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6028 as residues: Thr-6 to Gly-13, Trp-20 to Thr-36. 877092 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6029 as residues: Arg-17 to Gly-23. 877093 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6030 as residues: Pro-33 to Cys-43. 877094 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6031 as residues: Pro-9 to Tyr-17, Gln-29 to Tyr-38, Ala-47 to Glu-55. 877096 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6031 as residues: Pro-9 to Tyr-17, Gln-29 to Tyr-38, Ala-47 to Glu-55. 877096 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6031 as residues: Pro-9 to Tyr-17, Gln-29 to Tyr-38, Ala-47 to Glu-55.	877066	
Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6017 as residues: Pro-11 to Asp-16, Arg-23 to Gln-29. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6018 as residues: Lys-26 to Arg-32. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6020 as residues: Arg-33 to Leu-40. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6021 as residues: Pro-23 to Asn-31, Leu-33 to Phe-38. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6022 as residues: Ser-1 to Ser-17. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6027 as residues: Arg-1 to Met-6, Thr-34 to Glu-54, Glu-58 to Asn-63. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6028 as residues: Thr-6 to Gly-13, Trp-20 to Thr-36. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6029 as residues: Arg-17 to Gly-23. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6030 as residues: Pro-33 to Cys-43. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6031 as residues: Pro-3 to Tyr-17, Gln-29 to Tyr-38, Ala-47 to Glu-55. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6031 as residues: Pro-9 to Tyr-17, Gln-29 to Tyr-38, Ala-47 to Glu-55. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6033 as residues: Pro-9 to Tyr-17, Gln-29 to Tyr-38, Ala-47 to Glu-55.	877000	
ID NO. 6017 as residues: Pro-11 to Asp-16, Arg-23 to Gln-29. 877068 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6018 as residues: Lys-26 to Arg-32. 877070 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6020 as residues: Arg-33 to Leu-40. 877071 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6021 as residues: Pro-23 to Asn-31, Leu-33 to Phe-38. 877073 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6022 as residues: Ser-1 to Ser-17. 877087 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6027 as residues: Arg-1 to Met-6, Thr-34 to Glu-54, Glu-58 to Asn-63. 877088 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6028 as residues: Thr-6 to Gly-13, Trp-20 to Thr-36. 877092 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6029 as residues: Arg-17 to Gly-23. 877093 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6030 as residues: Pro-33 to Cys-43. 877094 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6031 as residues: Pro-9 to Tyr-17, Gln-29 to Tyr-38, Ala-47 to Glu-55. 877096 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6033 as residues: Pro-9 to Tyr-17, Gln-29 to Tyr-38, Ala-47 to Glu-55.	077047	
Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6018 as residues: Lys-26 to Arg-32. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6020 as residues: Arg-33 to Leu-40. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6021 as residues: Pro-23 to Asn-31, Leu-33 to Phe-38. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6022 as residues: Ser-1 to Ser-17. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6027 as residues: Arg-1 to Met-6, Thr-34 to Glu-54, Glu-58 to Asn-63. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6028 as residues: Thr-6 to Gly-13, Trp-20 to Thr-36. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6029 as residues: Arg-17 to Gly-23. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6030 as residues: Pro-33 to Cys-43. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6031 as residues: Pro-9 to Tyr-17, Gln-29 to Tyr-38, Ala-47 to Glu-55. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6033 as residues: Pro-9 to Tyr-17, Gln-29 to Tyr-38, Ala-47 to Glu-55. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6033 as residues: Lys-9 to Ser-17.	8//06/	
ID NO. 6018 as residues: Lys-26 to Arg-32. 877070 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6020 as residues: Arg-33 to Leu-40. 877071 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6021 as residues: Pro-23 to Asn-31, Leu-33 to Phe-38. 877073 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6022 as residues: Ser-1 to Ser-17. 877087 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6027 as residues: Arg-1 to Met-6, Thr-34 to Glu-54, Glu-58 to Asn-63. 877088 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6028 as residues: Thr-6 to Gly-13, Trp-20 to Thr-36. 877092 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6029 as residues: Arg-17 to Gly-23. 877093 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6030 as residues: Pro-33 to Cys-43. 877094 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6031 as residues: Pro-9 to Tyr-17, Gln-29 to Tyr-38, Ala-47 to Glu-55. 877096 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6033 as residues: Pro-9 to Tyr-17, Gln-29 to Tyr-38, Ala-47 to Glu-55.		
Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6020 as residues: Arg-33 to Leu-40. 877071 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6021 as residues: Pro-23 to Asn-31, Leu-33 to Phe-38. 877073 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6022 as residues: Ser-1 to Ser-17. 877087 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6027 as residues: Arg-1 to Met-6, Thr-34 to Glu-54, Glu-58 to Asn-63. 877088 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6028 as residues: Thr-6 to Gly-13, Trp-20 to Thr-36. 877092 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6029 as residues: Arg-17 to Gly-23. 877093 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6030 as residues: Pro-33 to Cys-43. 877094 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6031 as residues: Pro-9 to Tyr-17, Gln-29 to Tyr-38, Ala-47 to Glu-55. 877096 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6033 as residues: Pro-9 to Tyr-17, Gln-29 to Tyr-38, Ala-47 to Glu-55.	877068	
ID NO. 6020 as residues: Arg-33 to Leu-40. 877071 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6021 as residues: Pro-23 to Asn-31, Leu-33 to Phe-38. 877073 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6022 as residues: Ser-1 to Ser-17. 877087 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6027 as residues: Arg-1 to Met-6, Thr-34 to Glu-54, Glu-58 to Asn-63. 877088 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6028 as residues: Thr-6 to Gly-13, Trp-20 to Thr-36. 877092 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6029 as residues: Arg-17 to Gly-23. 877093 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6030 as residues: Pro-33 to Cys-43. 877094 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6031 as residues: Pro-9 to Tyr-17, Gln-29 to Tyr-38, Ala-47 to Glu-55. 877096 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6033 as residues: Pro-9 to Tyr-17, Gln-29 to Tyr-38, Ala-47 to Glu-55.		ID NO. 6018 as residues: Lys-26 to Arg-32.
ID NO. 6020 as residues: Arg-33 to Leu-40. 877071 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6021 as residues: Pro-23 to Asn-31, Leu-33 to Phe-38. 877073 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6022 as residues: Ser-1 to Ser-17. 877087 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6027 as residues: Arg-1 to Met-6, Thr-34 to Glu-54, Glu-58 to Asn-63. 877088 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6028 as residues: Thr-6 to Gly-13, Trp-20 to Thr-36. 877092 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6029 as residues: Arg-17 to Gly-23. 877093 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6030 as residues: Pro-33 to Cys-43. 877094 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6031 as residues: Pro-9 to Tyr-17, Gln-29 to Tyr-38, Ala-47 to Glu-55. 877096 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6033 as residues: Pro-9 to Tyr-17, Gln-29 to Tyr-38, Ala-47 to Glu-55.	877070	Preferred epitopes include those comprising a sequence shown in SEQ
Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6021 as residues: Pro-23 to Asn-31, Leu-33 to Phe-38. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6022 as residues: Ser-1 to Ser-17. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6027 as residues: Arg-1 to Met-6, Thr-34 to Glu-54, Glu-58 to Asn-63. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6028 as residues: Thr-6 to Gly-13, Trp-20 to Thr-36. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6029 as residues: Arg-17 to Gly-23. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6030 as residues: Pro-33 to Cys-43. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6031 as residues: Pro-9 to Tyr-17, Gln-29 to Tyr-38, Ala-47 to Glu-55. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6033 as residues: Pro-9 to Tyr-17, Gln-29 to Tyr-38, Ala-47 to Glu-55. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6033 as residues: Pro-9 to Tyr-17, Gln-29 to Tyr-38, Ala-47 to Glu-55.		
ID NO. 6021 as residues: Pro-23 to Asn-31, Leu-33 to Phe-38. 877073 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6022 as residues: Ser-1 to Ser-17. 877087 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6027 as residues: Arg-1 to Met-6, Thr-34 to Glu-54, Glu-58 to Asn-63. 877088 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6028 as residues: Thr-6 to Gly-13, Trp-20 to Thr-36. 877092 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6029 as residues: Arg-17 to Gly-23. 877093 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6030 as residues: Pro-33 to Cys-43. 877094 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6031 as residues: Pro-9 to Tyr-17, Gln-29 to Tyr-38, Ala-47 to Glu-55. 877096 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6033 as residues: Lys-9 to Ser-17.	877071	
Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6022 as residues: Ser-1 to Ser-17. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6027 as residues: Arg-1 to Met-6, Thr-34 to Glu-54, Glu-58 to Asn-63. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6028 as residues: Thr-6 to Gly-13, Trp-20 to Thr-36. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6029 as residues: Arg-17 to Gly-23. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6030 as residues: Pro-33 to Cys-43. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6031 as residues: Pro-9 to Tyr-17, Gln-29 to Tyr-38, Ala-47 to Glu-55. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6033 as residues: Pro-9 to Tyr-17, Gln-29 to Tyr-38, Ala-47 to Glu-55. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6033 as residues: Lys-9 to Ser-17.		
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Asn-63. 877088 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6028 as residues: Thr-6 to Gly-13, Trp-20 to Thr-36. 877092 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6029 as residues: Arg-17 to Gly-23. 877093 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6030 as residues: Pro-33 to Cys-43. 877094 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6031 as residues: Pro-9 to Tyr-17, Gln-29 to Tyr-38, Ala-47 to Glu-55. 877096 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6033 as residues: Lys-9 to Ser-17.	0//08/	
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Glu-55. 877096 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6033 as residues: Lys-9 to Ser-17.	377034	
877096 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6033 as residues: Lys-9 to Ser-17.		
ID NO. 6033 as residues: Lys-9 to Ser-17.	07500	
	877096	
877097 Preferred epitopes include those comprising a sequence shown in SEQ		
	877097	Preferred epitopes include those comprising a sequence shown in SEQ

	ID NO. 6034 as residues: Phe-34 to Ser-39, Glu-63 to Phe-74, Leu-78 to Pro-83.
877098	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6035 as residues: Lys-1 to Asp-8.
077000	
877099	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6036 as residues: Pro-10 to Gly-17, Tyr-23 to Ser-28.
877101	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6037 as residues: Asp-22 to Cys-28, Gly-87 to Leu-93, Lys-128
	to Asn-151.
877105	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6039 as residues: Pro-48 to Cys-53.
877106	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6040 as residues: Gln-3 to Ile-12.
877110	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6041 as residues: Val-6 to Ala-13.
877111	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6042 as residues: Phe-56 to Asn-72, Gln-84 to Leu-93, Ser-96
	to Pro-109, Pro-116 to Glu-126.
877114	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6044 as residues: Lys-13 to Lys-21.
877119	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6045 as residues: Ala-16 to Ser-22.
877120	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6046 as residues: Pro-1 to Gly-14, Gly-33 to Ser-40, Gln-80 to
	Ser-101.
877121	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6047 as residues: Arg-34 to Ser-40.
877123	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6049 as residues: Thr-33 to Asp-38.
877126	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6050 as residues: Gly-10 to Leu-22, Gly-47 to Lys-62.
877132	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6054 as residues: Ser-2 to Lys-8.
877133	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6055 as residues: Thr-1 to Asp-8.
877135	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6057 as residues: Leu-7 to Leu-13, Pro-15 to Cys-28, Ser-50 to
	Lys-56.
877137	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6058 as residues: Glu-65 to Arg-72.
877138	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6059 as residues: Lys-15 to Thr-21.
877140	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6061 as residues: Ile-45 to Phe-51.
877142	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6062 as residues: Thr-5 to Ser-12.
877143	Preferred epitopes include those comprising a sequence shown in SEQ
3,,,,,,,,,,	ID NO. 6063 as residues: Arg-1 to Leu-6.
877148	Preferred epitopes include those comprising a sequence shown in SEQ
0,,,,,	1

	ID NO. 6067 as residues: Leu-32 to Trp-37.
877149	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6068 as residues: Lys-72 to Gln-86.
877153	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6069 as residues: Cys-40 to Cys-46.
877154	Preferred epitopes include those comprising a sequence shown in SEQ
0,,,,,,	ID NO. 6070 as residues: Asn-24 to Phe-29, Thr-45 to Lys-50.
877165	Preferred epitopes include those comprising a sequence shown in SEQ
8//103	
077166	ID NO. 6074 as residues: Arg-6 to Lys-11, His-20 to Asn-25.
877166	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6075 as residues: Tyr-1 to Arg-7.
877167	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6076 as residues: Glu-25 to Asn-34.
877168	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6077 as residues: Tyr-1 to Ile-6, Val-17 to Ser-23, Thr-35 to
	His-40.
877169	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6078 as residues: Pro-1 to Met-12.
877170	Preferred epitopes include those comprising a sequence shown in SEQ
3,,,,,,	ID NO. 6079 as residues: Ser-4 to Lys-9.
877171	
0//1/1	Preferred epitopes include those comprising a sequence shown in SEQ
053153	ID NO. 6080 as residues: Val-10 to Leu-15, Arg-34 to Leu-40.
877173	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6081 as residues: Pro-18 to Gly-31.
877174	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6082 as residues: Lys-16 to Gln-21.
877175	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6083 as residues: Glu-2 to Ser-9.
877181	Preferred epitopes include those comprising a sequence shown in SEQ
V	ID NO. 6085 as residues: Glu-16 to Glu-23.
877187	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6088 as residues: Asp-41 to Ile-50, Thr-73 to Val-89, Gln-118
	to Asp-123.
877194	Preferred epitopes include those comprising a sequence shown in SEQ
0//1/4	
877195	ID NO. 6091 as residues: Gly-53 to Asp-63.
0//193	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6092 as residues: Pro-17 to Ile-24, Pro-28 to Phe-34.
877200	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6093 as residues: Thr-29 to Lys-35, Asp-44 to Cys-49.
877202	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6094 as residues: Gly-17 to Ala-23, Leu-52 to Asn-58.
877205	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6095 as residues: Lys-12 to Asp-18, Leu-40 to Arg-67, Val-75
	to Asp-84.
877207	Preferred epitopes include those comprising a sequence shown in SEQ
[0,720,	ID NO. 6097 as residues: Ala-19 to Arg-29.
877208	
0//208	Preferred epitopes include those comprising a sequence shown in SEQ
0770::	ID NO. 6098 as residues: Tyr-4 to Gln-9.
877211	Preferred epitopes include those comprising a sequence shown in SEQ

	ID NO. 6099 as residues: Asp-12 to Arg-17, Asp-34 to Gln-43, Asn-78
	to Glu-84, Ser-99 to Ala-105, Ser-108 to His-113, Ile-115 to Gly-122,
	Phe-132 to Arg-148.
877212	Preferred epitopes include those comprising a sequence shown in SEQ
07,7212	ID NO. 6100 as residues: Gln-1 to Ser-9.
877213	Preferred epitopes include those comprising a sequence shown in SEQ
677213	ID NO. 6101 as residues: Arg-42 to Gln-53, His-56 to Ala-62, Asn-73
	to Pro-81.
077014	
877214	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6102 as residues: Ser-15 to Cys-23.
877218	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6104 as residues: Lys-33 to Phe-40, Pro-64 to Arg-72, Arg-105
	to Gly-110.
877220	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6105 as residues: Gly-1 to Thr-14, Ala-27 to Leu-32, Pro-47 to
	Pro-54.
877230	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6108 as residues: Thr-1 to Asn-8.
877231	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6109 as residues: Gly-1 to Ser-20, Phe-29 to Asn-37, Asn-55 to
	Tyr-64, Ala-69 to Asp-78, Tyr-82 to Ala-91, Lys-100 to Glu-122.
877232	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6110 as residues: Lys-41 to Ile-47.
877233	Preferred epitopes include those comprising a sequence shown in SEQ
, ,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,	ID NO. 6111 as residues: Ile-11 to Phe-16, Tyr-27 to Pro-33.
877234	Preferred epitopes include those comprising a sequence shown in SEQ
077254	ID NO. 6112 as residues: Ala-13 to His-18, Gly-24 to Thr-29, Pro-31 to
	Gly-39, Pro-49 to Asp-56, Trp-64 to Asp-72, Pro-74 to Asp-80.
877235	Preferred epitopes include those comprising a sequence shown in SEQ
677233	ID NO. 6113 as residues: Thr-6 to Gly-12, Pro-41 to Asp-48, Gly-54 to
	1
	Phe-62, His-94 to Tyr-102, Ser-108 to Gly-123, Gln-130 to Asn-136,
027240	Tyr-169 to His-175, Phe-188 to Arg-195, Trp-232 to Ile-237.
877240	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6115 as residues: His-1 to Leu-8, Ala-42 to Arg-50, His-74 to
	Tyr-81.
877242	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6116 as residues: Asp-25 to Asn-30.
877250	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6118 as residues: Thr-11 to Cys-22, Gly-29 to Gly-37, Arg-74
	to Asn-91, Phe-110 to Pro-119, Thr-144 to Gln-149, Tyr-165 to Gly-
	171, Pro-190 to Ser-196.
877251	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6119 as residues: Ala-5 to Ser-11, Thr-32 to Thr-37, Gln-46 to
	Asp-57, Ala-70 to Gly-78.
877254	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6120 as residues: Val-50 to Tyr-55, Thr-63 to Thr-68, Phe-77 to
	Gly-92, Arg-112 to Lys-119.
877258	Preferred epitopes include those comprising a sequence shown in SEQ
3.,255	ID NO. 6124 as residues: Thr-1 to Ser-6, Thr-40 to Trp-49, Asn-65 to
	Lys-72.
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877263	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6126 as residues: Asp-1 to Ser-16, Pro-21 to Glu-26, Pro-46 to
877264	Asn-55, Thr-74 to Leu-86, Ser-96 to Asp-105.
877204	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6127 as residues: Thr-1 to Arg-6, Ser-14 to Arg-20.
877272	Preferred epitopes include those comprising a sequence shown in SEQ
077272	ID NO. 6128 as residues: Ile-55 to Leu-69, Thr-84 to Pro-94, Pro-104 to
	His-120.
877274	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6129 as residues: Glu-50 to Pro-58, Ile-88 to Gly-97, Pro-107 to
	Gly-116, Gln-136 to Gly-142, Asp-164 to Glu-176.
877275	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6130 as residues: Pro-1 to Gln-19, Cys-27 to Thr-34, Ile-49 to
077201	Trp-56.
877281	Preferred epitopes include those comprising a sequence shown in SEQ
877282	ID NO. 6132 as residues: Lys-17 to Thr-23.
0//202	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6133 as residues: Ala-1 to Lys-7, Asp-12 to Phe-17, Ile-24 to Glu-43.
877283	Preferred epitopes include those comprising a sequence shown in SEQ
0,7203	ID NO. 6134 as residues: Lys-18 to Ile-23.
877284	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6135 as residues: Ile-41 to Trp-46, Glu-64 to Gly-80, Glu-134
	to Gly-141, Phe-143 to Ser-158, Gln-207 to Asp-212.
877285	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6136 as residues: His-1 to Leu-11.
877290	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6140 as residues: Pro-34 to Tyr-40, Ser-67 to Trp-73, Asp-103
	to Phe-109, Gln-130 to Gly-135, Trp-188 to Trp-197, Leu-327 to Asn-
	333, Gly-401 to Asn-407, Asn-473 to Val-483, Ser-523 to Gln-529, Arg-
	538 to Ser-544, Ala-563 to Ser-573, Gln-581 to Thr-592.
877295	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6141 as residues: Gln-54 to Leu-66, Pro-74 to Asp-79, Val-104
	to Leu-112, Asn-114 to Asn-122, Glu-141 to Lys-152, Pro-265 to Leu-
	271, Phe-275 to Ser-280, Glu-298 to Ala-304, Arg-317 to Leu-323, Gln-
877298	332 to Tyr-337, Gln-342 to Arg-352.
07/290	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6142 as residues: Ser-60 to Gly-66.
877301	Preferred epitopes include those comprising a sequence shown in SEQ
] 377301	ID NO. 6144 as residues: Gln-17 to Lys-24, Ala-28 to Cys-35.
877310	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6145 as residues: Met-2 to Leu-12, Ser-16 to Asp-23, Gly-38 to
	Lys-45.
877319	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6146 as residues: Ala-30 to Glu-44.
877321	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6148 as residues: Gln-1 to Arg-7.
877326	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6150 as residues: Thr-25 to Asp-31.

877327	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6151 as residues: Thr-3 to Ser-10.
877332	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6154 as residues: Gly-26 to Arg-43.
877333	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6155 as residues: Pro-10 to Trp-19.
877334	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6156 as residues: Ala-18 to Ala-32, Thr-52 to Ser-60.
877336	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6157 as residues: Cys-10 to Phe-17.
877340	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6160 as residues: Ser-32 to Arg-38, Ala-72 to Lys-79, Arg-103 to Phe-111.
877344	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6161 as residues: His-41 to Thr-48.
877346	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6162 as residues: Ala-66 to Gln-78.
877355	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6165 as residues: Ser-12 to His-21, Pro-59 to Asp-69.
877356	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6166 as residues: Ser-12 to His-21, Pro-59 to Glu-68.
877361	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6168 as residues: Pro-1 to Ser-7, Thr-45 to Leu-63, Arg-113 to Thr-118, Pro-172 to Gly-182.
877370	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6170 as residues: Asp-17 to Gly-23, Lys-89 to Asp-94, Lys-129 to Asp-134, Leu-195 to Glu-204, Asn-325 to Val-336.
877373	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6171 as residues: His-8 to Gly-18, Gln-56 to Arg-61, Arg-160 to Pro-170, Ala-200 to Ser-212, His-225 to Lys-231, Gly-245 to Lys-254, Tyr-257 to Tyr-277, Pro-279 to Thr-287, Pro-305 to Gly-327, Tyr-342 to Glu-348.
877375	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6172 as residues: Gln-1 to Ser-22, Lys-40 to Phe-48, Leu-52 to His-57.
877377	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6173 as residues: Ser-27 to Thr-42, Lys-71 to Lys-85, Gly-99 to Arg-105.
877378	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6174 as residues: Lys-25 to Lys-39, Gly-53 to Arg-59, Ser-172 to Val-181.
877380	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6175 as residues: Glu-7 to Arg-20, Thr-28 to Trp-44, Ser-110 to Lys-118, Pro-124 to Arg-130, Ala-137 to Asn-147.
877384	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6176 as residues: Thr-2 to Leu-9, Thr-12 to Gly-17, Glu-26 to Ser-61, Asn-70 to Cys-80, Cys-84 to Ala-91, Lys-111 to Ser-119, Asn-170 to Gln-183, Ser-203 to Lys-210, Gln-216 to Pro-229, Arg-238 to Trp-255, Ile-257 to Phe-269.

877387	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6177 as residues: Asp-12 to Tyr-18, Pro-57 to Leu-63, Glu-90
	to Ala-96, Gly-102 to Val-111, Gln-123 to Ile-129, Asp-143 to Ala-150,
	Lys-156 to Arg-161, Thr-213 to Cys-220, Arg-256 to Tyr-261, Ser-265
	to Asp-274, Asp-290 to Ser-297, Val-307 to Arg-313, Asp-324 to Lys-
	337, Ser-438 to Arg-443, Asn-580 to Glu-585.
877388	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6178 as residues: Gly-15 to Asn-22.
877390	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6179 as residues: Cys-7 to Gly-24, Thr-31 to Val-53, Trp-102
	to Glu-108, Thr-118 to Gly-124.
877393	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6180 as residues: Glu-4 to Trp-9.
877406	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6182 as residues: Asn-1 to Glu-27, Lys-37 to Lys-46, Arg-59 to
	Lys-83, Asn-89 to Phe-95, His-102 to Asn-107, Ser-155 to Ile-168, Pro-
	175 to Gln-188, Asn-201 to Pro-211, Ala-234 to Ile-239, Asn-249 to
	Val-257, Pro-261 to Gly-275.
877408	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6183 as residues: Gln-1 to Pro-16, Pro-21 to Pro-30, Gly-47 to
	Gly-65, Tyr-78 to Leu-86, Glu-88 to Pro-104, Glu-118 to Ala-131, Ala-
	143 to Trp-150, Asp-152 to Ser-157, Ser-180 to Trp-187, Ser-190 to
	Pro-197, Ala-211 to Asn-219, Asp-252 to Leu-257, Thr-287 to Val-295.
877411.	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6184 as residues: His-20 to Gln-25, Asn-36 to Ser-56.
877630	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6186 as residues: Gln-40 to Phe-45.
878274	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6191 as residues: Pro-6 to Trp-14, Tyr-19 to Leu-26, Pro-56 to
	His-66, Tyr-70 to Arg-80, Thr-83 to Leu-100, Cys-107 to Phe-112, Lys-
	137 to Arg-148, Pro-155 to Leu-162.
878374	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6192 as residues: Arg-20 to Leu-28, Phe-57 to Arg-79.
878403	Preferred epitopes include those comprising a sequence shown in SEQ
`	ID NO. 6193 as residues: Ser-2 to Thr-8.
878433	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6194 as residues: Asn-17 to His-24, Pro-97 to Glu-111.
878436	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6195 as residues: Ser-18 to Thr-25.
878560	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6196 as residues: Thr-33 to Pro-40, Asp-62 to Glu-67, Ser-104
	to Phe-109.
878800	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6197 as residues: Leu-24 to Arg-30.
878909	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6198 as residues: Pro-14 to Ser-19, His-40 to Trp-49.
878917	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6199 as residues: Glu-26 to Thr-32, Ser-41 to Pro-46, Leu-107
	to Glu-115.
879009	Preferred epitopes include those comprising a sequence shown in SEQ

	ID NO. 6201 as residues: Trp-60 to His-68, Pro-99 to Gly-106.
879234	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6202 as residues: Ser-46 to Thr-64, Thr-69 to Gly-79, Ser-102
	to Arg-115, Leu-137 to Thr-144, Ala-146 to Pro-153, Pro-163 to Arg-
	180, Cys-209 to His-229.
879386	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6203 as residues: Pro-3 to Cys-11, Pro-70 to Phe-83, Ser-101 to
	Leu-107, Glu-110 to Pro-116, Lys-153 to Arg-158.
879484	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6204 as residues: Lys-44 to His-50, Thr-110 to Pro-116, Lys-
	178 to Gln-183, Pro-196 to Lys-205, Arg-214 to Thr-220, Asp-295 to
	Leu-301, Pro-316 to Glu-324, Glu-331 to Tyr-336, Gly-347 to Val-354.
879595	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6205 as residues: Pro-7 to Ser-15, Gly-49 to Ala-55, Gln-74 to
1	Pro-86.
879661	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6206 as residues: Arg-10 to Arg-20, Gly-48 to Val-53, Glu-69
	to Asp-76, Glu-116 to Glu-122, Glu-132 to Trp-143, Asp-166 to Asn-
	175, Arg-191 to Asn-197, Gln-205 to Gly-233, Lys-235 to Ala-274.
880071	Preferred epitopes include those comprising a sequence shown in SEQ
000071	ID NO. 6208 as residues: Ser-36 to Ser-41, Ser-77 to Gln-83.
880074	Preferred epitopes include those comprising a sequence shown in SEQ
000074	ID NO. 6209 as residues: Ser-7 to Gln-12, Gly-25 to Gly-31, Gly-71 to
	Gly-84, Leu-147 to Glu-164, Trp-172 to Leu-180.
880418	Preferred epitopes include those comprising a sequence shown in SEQ
000410	ID NO. 6210 as residues: Ser-56 to Val-64, Lys-66 to Cys-73.
880649	Preferred epitopes include those comprising a sequence shown in SEQ
000042	ID NO. 6212 as residues: His-28 to Gly-35, Gln-141 to His-147, Glu-
	232 to Gln-237, Ala-264 to Glu-269.
880694	Preferred epitopes include those comprising a sequence shown in SEQ
000074	ID NO. 6213 as residues: Glu-21 to Glu-27, Arg-34 to Ile-41, Leu-83 to
	Ala-93, Pro-120 to Glu-130.
880747	Preferred epitopes include those comprising a sequence shown in SEQ
000747	ID NO. 6214 as residues: Pro-16 to Phe-23, Gln-45 to Cys-50, Asn-66
	to Asn-73, Ile-98 to His-105, Pro-183 to Pro-190, His-206 to Ser-212,
Į.	Thr-295 to Pro-316, Ser-364 to Trp-370, Gln-385 to Asn-396.
880994	Preferred epitopes include those comprising a sequence shown in SEQ
000974	ID NO. 6216 as residues: Ile-32 to Tyr-47.
881105	Preferred epitopes include those comprising a sequence shown in SEQ
001103	ID NO. 6220 as residues: Arg-9 to Gln-35, Ile-113 to Gly-120.
001210	Preferred epitopes include those comprising a sequence shown in SEQ
881219	ID NO. 6221 as residues: Ser-17 to Thr-25, Lys-39 to Thr-48, His-53 to
	Arg-60, Pro-67 to Asn-72, Thr-157 to Phe-165, Gln-212 to Glu-221, Gly-241 to Ser-260, Thr-294 to Phe-300, Ile-319 to Lys-328, Ser-338 to
	Lys-343, Leu-383 to Phe-388, Gly-430 to Asp-441, Ser-466 to Glu-475,
	Gln-541 to Pro-554, Val-583 to Thr-595, Leu-598 to Arg-603, Gln-608
	to Gln-614, Asp-639 to Asn-648, Asp-654 to Phe-667, Lys-676 to Val-
001221	704, Lys-725 to Ser-731, Pro-739 to Ala-763, Asp-772 to Gly-778.
881221	Preferred epitopes include those comprising a sequence shown in SEQ
L	ID NO. 6222 as residues: Ile-1 to Lys-11, Asn-59 to Phe-65, Phe-70 to

	Asn-79, Lys-156 to Glu-162, Pro-168 to Asp-175, Pro-213 to Leu-219,
	Asn-246 to Leu-266, Ser-275 to Asp-286, Gln-334 to Leu-345.
882330	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6223 as residues: Arg-20 to Ser-27, Glu-40 to Glu-50.
882715	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6224 as residues: Glu-4 to Asn-14, Gln-66 to Gly-73, Leu-88 to
	Leu-97, Val-101 to Gln-107.
882729	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6225 as residues: Arg-7 to Gly-12, Met-42 to Ser-58, Gln-65 to
	Asn-73, Glu-91 to Ala-99, Pro-103 to Tyr-109, Arg-174 to Ala-179, His-
	189 to Gln-196, Asn-208 to Pro-219.
882762	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6226 as residues: Arg-8 to Asn-30, Ser-37 to Gln-42, His-74 to
200150	Leu-82, Arg-92 to His-97, Gln-114 to Leu-119, Gly-131 to Gly-137.
883172	Preferred epitopes include those comprising a sequence shown in SEQ
002271	ID NO. 6227 as residues: His-1 to Arg-10.
883371	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6230 as residues: Asp-24 to Trp-41, Tyr-106 to Lys-114, Ala-
	161 to Glu-167, Pro-182 to Leu-190, Ala-193 to Pro-200, Leu-205 to
883753	Tyr-212, Pro-240 to Lys-252, Pro-254 to Lys-262, Leu-293 to Leu-303. Preferred epitopes include those comprising a sequence shown in SEQ
003733	ID NO. 6231 as residues: Gly-156 to Met-161, Cys-186 to Lys-197.
883799	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6232 as residues: Ser-1 to Glu-18, Val-79 to Glu-88.
883945	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6233 as residues: Ser-21 to Arg-28.
883971	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6234 as residues: Ser-19 to Gly-24, Gly-54 to Ser-59.
884038	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6235 as residues: Pro-18 to Asn-25, Ala-44 to Asn-50, Arg-56
	to Lys-64, Gly-76 to Gly-85, Lys-92 to Leu-98, Gly-116 to Gly-121,
004006	Gln-132 to His-138, Thr-159 to Asp-167.
884095	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6236 as residues: Arg-50 to Thr-56, Pro-116 to Arg-121, Lys-
884161	129 to Phe-136, Glu-139 to Leu-144, Lys-156 to Leu-162.
884101	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6237 as residues: Asn-16 to Tyr-23, Glu-47 to Trp-56, Ser-90 to Lys-96, Ala-126 to Glu-136, Pro-138 to Lys-149, Glu-181 to Gly-186,
	Trp-208 to Lys-219, Arg-347 to Ala-358, Leu-370 to Lys-381, Thr-408
	to Ile-415, Pro-425 to Leu-437, Gln-450 to Asn-455.
884168	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6238 as residues: Glu-94 to Tyr-102, Pro-105 to Asn-112, Thr-
	121 to Gly-137, Glu-157 to Gly-162, Glu-179 to Phe-186, Cys-211 to
	Thr-222, Ser-240 to Lys-245, Thr-262 to Asn-279, Arg-288 to Pro-306,
	Asn-332 to Gln-339, Ser-375 to Leu-382, Arg-408 to Gly-415, Asp-423
t	to Thr-428, Ser-471 to Asn-476, Pro-545 to Gly-551, Ser-606 to Pro-
	616, Ala-662 to Gly-667, Thr-675 to Tyr-682, Glu-714 to Trp-720, Pro-
	722 to Val-732, Pro-787 to Thr-795, Arg-811 to Glu-816, Gln-880 to
	Thr-891.
884215	Preferred epitopes include those comprising a sequence shown in SEQ

	ID NO. 6239 as residues: Met-10 to Gln-18, Pro-23 to Leu-31, Glu-46
	to Arg-51, Phe-135 to Pro-143, His-218 to Asp-227, Pro-244 to Met-
	250, Lys-258 to Asp-263, Pro-266 to Leu-276, Pro-286 to Asp-293.
884529	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6241 as residues: Arg-8 to Ser-15, Gln-89 to Gln-95, Leu-109
	to Tyr-115, Glu-126 to Arg-133.
884719	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6242 as residues: Arg-4 to Ala-10, Arg-40 to Gly-45, Asp-86 to
	Tyr-91, Pro-100 to Phe-113.
885350	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6243 as residues: Arg-15 to Pro-21, Cys-29 to Cys-41, Pro-52
	to Leu-63, Pro-98 to Ser-108, Tyr-113 to Cys-118, Cys-124 to Asp-129,
	Cys-180 to Gln-187, Cys-247 to Cys-259, Ser-279 to Trp-286, Cys-296
	to Cys-302, Pro-304 to Cys-309, Ser-343 to His-348, Gln-367 to Lys-
	373.
885476	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6244 as residues: Lys-28 to Glu-51, Lys-123 to Leu-133.
885484	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6245 as residues: Arg-1 to Glu-10, Gly-22 to Gly-27.
886505	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6248 as residues: Ser-64 to Gln-70, Ala-75 to Leu-80, His-82 to
	Gly-87, Ser-121 to Lys-137.
886788	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6250 as residues: Lys-60 to Lys-65, Lys-78 to Lys-94, Leu-116
	to Gln-123.
887098	Preferred epitopes include those comprising a sequence shown in SEQ
1	ID NO. 6252 as residues: Pro-1 to Ala-9, Val-56 to Val-63, Gly-86 to
	Glu-91.
887114	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6253 as residues: Glu-38 to Arg-52, Ser-56 to Val-62.
887155	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6254 as residues: Thr-3 to Pro-9, Pro-18 to Gly-25, Ala-30 to
	Gly-36, Arg-41 to Asp-56, Ala-60 to Pro-68, Met-99 to Leu-128, Thr-
	143 to Phe-157.
887172	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6255 as residues: Cys-5 to Ser-14, Val-83 to Ser-88.
887192	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6256 as residues: Glu-29 to Cys-39, Val-46 to Ser-52, Asn-58
	to Gly-65, Cys-68 to His-82, Tyr-84 to Gly-94, Leu-122 to Trp-138,
	Ala-158 to Leu-170, Gly-175 to Arg-182, Tyr-203 to Ser-210, Gly-246
	to Met-258, Arg-288 to Gln-296.
887280	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6257 as residues: Asn-1 to Gly-15, Pro-18 to Asn-28, Gln-35 to
005000	Glu-40, Arg-60 to Arg-69.
887399	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6258 as residues: Pro-8 to Gly-18, Ala-94 to Gly-99, Asn-107
	to Arg-112, Phe-161 to Arg-166, Thr-196 to Phe-201, Tyr-309 to Gly-
00000	316, Leu-326 to Arg-331.
887535	Preferred epitopes include those comprising a sequence shown in SEQ
1	ID NO. 6261 as residues: Glu-26 to Gly-32, His-73 to Arg-79.

887803	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6262 as residues: Ala-1 to Gln-7, Lys-24 to Ser-30, Pro-44 to Ser-49, Ser-99 to Ser-105.
887857	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6263 as residues: Pro-1 to Ser-6, Pro-25 to Cys-31, Arg-142 to Lys-150, Pro-223 to Gly-230, Ala-233 to Val-247.
887892	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6264 as residues: Ser-10 to Ile-15, Val-60 to Arg-66, Tyr-114 to Leu-128.
887936	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6265 as residues: Leu-1 to Cys-6, Lys-46 to Thr-53, Ala-56 to Glu-63.
887996	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6266 as residues: Ala-1 to Gly-6, Pro-9 to Pro-24, Gln-70 to Tyr-82, Glu-127 to Ser-134.
888051	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6268 as residues: Trp-45 to Trp-56, Thr-58 to Asp-73, Thr-126 to Arg-133, Phe-148 to Ser-155, Val-208 to Gly-223.
888153	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6270 as residues: Gly-5 to Leu-12, Tyr-18 to Asp-25, Ile-88 to Ala-125, Ser-129 to Tyr-141, Gln-191 to Gln-196, Thr-290 to Asn-296, Thr-301 to Thr-309, Leu-360 to Ala-365, Leu-367 to Gly-378, Pro-398 to Gly-418, Pro-443 to Gly-454.
888402	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6272 as residues: Leu-11 to Asn-16, Gly-164 to Glu-171, Leu-181 to Ser-186, Asp-193 to Ser-201, Glu-222 to Leu-229, Gln-238 to Tyr-245, Leu-256 to Asp-267, Gly-286 to Gln-301, Ser-311 to Ala-319, Glu-345 to Gly-351, Phe-361 to Asp-367, Thr-436 to Arg-443, Ile-460 to Gln-467, Gln-510 to Glu-533, Ala-541 to Ala-548, Gln-561 to Glu-571, Leu-581 to Ala-590, Phe-639 to Ser-652.
888708	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6275 as residues: Ile-27 to Val-33, Ala-63 to Ser-69, Pro-128 to Ser-135.
888720	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6276 as residues: Phe-34 to Glu-44, Glu-111 to Gly-122.
888950	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6278 as residues: Lys-56 to Gln-64, Pro-172 to Gly-183, Asp-208 to Asn-216, Glu-227 to Gly-232, Pro-259 to Arg-269, Asn-281 to His-286.
889136 	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6279 as residues: Arg-1 to Lys-14, Glu-19 to His-26.
889263	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6280 as residues: Gly-18 to Gly-30.
889299	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6281 as residues: Leu-5 to Ser-12.
889300	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6282 as residues: Glu-15 to Gly-22, Asn-45 to Pro-51, Glu-141 to Asn-146, Asp-154 to Gln-163, Glu-185 to Ser-191, Arg-200 to Pro-206, Asp-220 to Asn-225, Glu-231 to Asn-237, Ser-262 to Gly-269, Pro-276 to Ala-281, Glu-314 to Thr-320, Ser-416 to His-424, Gly-426 to

	Ala-438, Pro-445 to Phe-450, Arg-464 to Leu-469.
889323	Preferred epitopes include those comprising a sequence shown in SEQ
007525	ID NO. 6283 as residues: Pro-1 to Gly-11, Pro-13 to His-42, Arg-55 to
	Arg-66, Arg-84 to Gly-91, Gly-96 to Pro-101, His-112 to Pro-118.
889368	Preferred epitopes include those comprising a sequence shown in SEQ
007500	ID NO. 6284 as residues: Pro-1 to Asn-9.
889467	Preferred epitopes include those comprising a sequence shown in SEQ
007407	ID NO. 6285 as residues: Asp-10 to Asp-19, Ala-63 to Asp-68.
889494	Preferred epitopes include those comprising a sequence shown in SEQ
005151	ID NO. 6286 as residues: Arg-1 to Ser-6.
889700	Preferred epitopes include those comprising a sequence shown in SEQ
002700	ID NO. 6287 as residues: Ala-4 to Gly-14, Pro-20 to Cys-27, Leu-88 to
	Gly-94, Gly-106 to Lys-120, Pro-144 to Leu-150.
889782	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6288 as residues: Val-103 to Ser-108.
889954	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6289 as residues: Glu-21 to Tyr-33, Ile-90 to Ser-95, Pro-103 to
	Val-111, Ala-133 to His-140, Asn-153 to Trp-159, Gln-187 to Glu-192,
	Lys-214 to Arg-224.
889994	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6291 as residues: Ala-1 to Gln-7, Lys-24 to Ser-30, Pro-44 to
	Ser-49.
890666	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6292 as residues: Pro-36 to Trp-51, Arg-96 to Gly-104, Glu-134
Ţ	to Asn-144, Pro-203 to His-210, Cys-228 to Asp-235, Gly-278 to Tyr-
	284, Ser-309 to Pro-316, Thr-325 to Ala-333, Ser-337 to Glu-357, Tyr-
	390 to Gly-403, Tyr-409 to Gly-421.
890698	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6293 as residues: Ser-37 to Asp-43.
890776	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6296 as residues: Ser-4 to Trp-13, Pro-276 to Ala-282, Ala-341
	to Arg-347.
890801	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6297 as residues: Asn-9 to Arg-15.
890820	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6298 as residues: Arg-110 to Asp-115, Leu-185 to Gln-193,
	Ser-201 to Asp-208, Arg-215 to Arg-221, Arg-242 to Tyr-250, Thr-315
	to Thr-320, Lys-359 to Val-367, Ser-395 to Tyr-401, Met-406 to Lys-
001064	411.
891264	Preferred epitopes include those comprising a sequence shown in SEQ
001205	ID NO. 6302 as residues: Asp-1 to Gly-15, Ala-22 to Tyr-28.
891305	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6303 as residues: Asp-39 to Tyr-44, Thr-46 to Asn-55, Ser-78
902112	to Ala-87. Preferred epitopes include those comprising a sequence shown in SEQ
892113	
902177	ID NO. 6305 as residues: Gln-15 to Gln-22, Leu-216 to Lys-223.
892177	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6306 as residues: His-8 to Gly-18, Glu-100 to Asn-107, Glu-
	121 to Asn-126, Lys-128 to Ala-140, Ala-180 to Arg-186, Phe-230 to
	Thr-238, Pro-325 to His-341.

892367	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6308 as residues: Ser-31 to Gln-40.
892563	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6310 as residues: Arg-1 to Gly-23.
892820	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6311 as residues: Pro-8 to Thr-19.
893457	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6313 as residues: Lys-12 to Thr-18.
893827	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6314 as residues: Glu-37 to Asn-42, Ser-48 to Thr-54, Pro-101 to Glu-106.
893842	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6315 as residues: Asp-1 to Tyr-7, His-71 to Pro-78.
893866	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6316 as residues: Ala-12 to Lys-28, Ala-88 to Gly-95, Thr-100 to Cys-109.
894012	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6318 as residues: Ser-39 to Gln-48, Ala-61 to Pro-69.
894051	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6319 as residues: Arg-52 to Glu-66.
894121	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6320 as residues: Gly-28 to Ser-36, Trp-38 to Pro-60, Pro-98 to Thr-104, Pro-113 to Tyr-118, Phe-133 to Gly-140, Pro-186 to Leu-192, Glu-239 to Gly-246, Pro-257 to Lys-269, Lys-273 to Lys-279.
894341	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6321 as residues: Asn-18 to Asp-29.
894631	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6323 as residues: Met-1 to Gly-17, Pro-22 to Gly-30, Gly-72 to His-82, Leu-89 to Lys-95.
894806	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6324 as residues: Leu-99 to Ser-104.
894811	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6325 as residues: Asn-1 to Asn-8, Phe-49 to Asn-54, Glu-57 to Ser-63.
894820	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6327 as residues: Leu-8 to Gly-15.
894824	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6328 as residues: Ser-8 to Asp-13, Arg-19 to Ser-25.
894827	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6329 as residues: Arg-5 to Lys-11.
894830	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6330 as residues: Thr-102 to Gln-132.
894831	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6331 as residues: Ile-132 to Gly-138, Phe-149 to Thr-154.
894832	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6332 as residues: Pro-6 to Lys-17, Ser-66 to Pro-72, Pro-84 to Val-93.
894842	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6333 as residues: Ser-65 to Asp-70.

894878	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6334 as residues: Arg-9 to Trp-27, Pro-39 to Asn-44.
895122	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6335 as residues: Thr-11 to Pro-34, Asn-151 to Glu-157, Asp-302 to Phe-309, Tyr-333 to Gly-339.
895303	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6336 as residues: His-1 to Asp-9, Leu-11 to Glu-24, Pro-59 to Gln-65.
895372	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6337 as residues: Asn-7 to Ser-19, Arg-81 to Asn-94, Lys-99 to Asp-104.
895675	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6338 as residues: Asn-47 to Gly-52, Pro-67 to Asp-72, Pro-100 to Leu-105, Ser-115 to Asp-120, Leu-128 to Asn-135.
895927	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6340 as residues: Asn-3 to Trp-18, Gly-30 to Ser-35, Pro-41 to Ser-51, Ser-132 to Tyr-143.
896008	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6341 as residues: Pro-5 to Thr-28, Val-65 to Gly-71, Thr-82 to Gly-96.
897234	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6342 as residues: Ala-1 to Asp-10, Leu-24 to Phe-30, Pro-36 to Ser-42.
897524	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6343 as residues: Thr-1 to Cys-24, Lys-26 to Ser-32, Gln-83 to Thr-91, Thr-131 to Gly-137, Lys-170 to Asp-177, Asp-190 to Pro-198.
897898	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6344 as residues: Pro-23 to Arg-31, Gln-79 to Gln-85, Cys-93 to Cys-107.
898087	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6345 as residues: Ser-49 to Asp-59, Arg-69 to Tyr-87.
898136	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6346 as residues: Ser-12 to Ser-19, Ala-47 to Lys-52, Arg-96 to His-105.
898192	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6348 as residues: His-9 to Ile-14, Tyr-58 to Phe-64, Thr-75 to Phe-81.
898355	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6349 as residues: Pro-5 to Gly-18, Pro-21 to Asn-31, Gln-38 to Glu-43, Arg-63 to Arg-78.
898427	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6351 as residues: Gly-6 to Ile-11, Pro-13 to Arg-38, Glu-68 to Lys-74, Asp-88 to Ser-93, Glu-122 to Gly-130, Glu-145 to Glu-150, Thr-156 to Asp-174, Glu-200 to Arg-208, Ala-226 to Leu-240.
898541	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6352 as residues: His-1 to Leu-11, Arg-37 to Ile-43, Gln-111 to Pro-120, Asp-133 to Asn-138, Arg-159 to Cys-165, Val-241 to Lys-265, Glu-326 to Tyr-331, Pro-365 to Asn-382, Asn-418 to Asp-430, Ala-434 to Ser-441, Tyr-479 to Gly-496, Pro-498 to Ser-505.
898651	Preferred epitopes include those comprising a sequence shown in SEQ

	ID NO. 6353 as residues: Ser-6 to Pro-11, Pro-27 to Glu-32, Pro-65 to Trp-71, Val-208 to Pro-215, His-220 to Thr-225.
898946	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6355 as residues: Thr-4 to Arg-14, Glu-34 to Pro-46.
899130	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6356 as residues: Pro-10 to His-19, Leu-47 to Tyr-55, Phe-93 to
ì	Gly-105, Ser-220 to Trp-227, Phe-295 to Thr-301, Thr-309 to Trp-315,
	Arg-326 to Phe-334, Arg-458 to Pro-466.
899224	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6357 as residues: Ser-3 to Gly-28, Gly-46 to Pro-56, Gly-70 to
	Ile-92, Gln-102 to Ser-117, Ala-123 to Pro-129, Pro-135 to Leu-140,
	Pro-150 to Asp-158, Pro-165 to Pro-177, Gln-188 to Asp-205, Ile-230 to
	Arg-245, His-251 to Trp-260, Asp-262 to Cys-267, Asn-296 to Arg-307,
	Glu-322 to Pro-330, Ile-351 to Asn-357, Asp-363 to Leu-369, Glu-386
000.000	to Phe-391, Lys-415 to Ser-420.
899632	Preferred epitopes include those comprising a sequence shown in SEQ
000664	ID NO. 6358 as residues: Thr-11 to Ser-16, Gly-25 to Asn-40.
899661	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6360 as residues: His-8 to Gly-18, Pro-35 to Trp-41, Arg-51 to
	Asp-64, Asp-69 to Gln-74, Gly-83 to Asn-96, Pro-107 to Lys-116, Glu-
200776	149 to Ser-171, Ile-177 to Ile-186.
899776	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6361 as residues: Met-36 to Arg-49, Pro-72 to Gly-82, Glu-89
200205	to Gly-96, Tyr-129 to Thr-135.
899885	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6363 as residues: His-65 to Gly-74, Asp-85 to Ser-97, Leu-133 to Glu-138, Glu-144 to Asp-153, Arg-170 to Ser-175, Gly-184 to Arg-
i	189, Gln-202 to Tyr-208.
899913	Preferred epitopes include those comprising a sequence shown in SEQ
0,,,,,	ID NO. 6364 as residues: Lys-1 to Tyr-16.
900015	Preferred epitopes include those comprising a sequence shown in SEQ
700015	ID NO. 6365 as residues: Lys-23 to His-36, Asp-52 to Leu-68.
900162	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6366 as residues: Gly-1 to Leu-9, Gly-48 to Gln-53, Cys-74 to
	Pro-79, Thr-118 to Val-128.
900555	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6368 as residues: His-8 to Gly-18, Cys-131 to Gly-136, Thr-198
	to Asn-203, Pro-231 to Asp-236.
900696	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6369 as residues: Arg-11 to Ser-23, Arg-72 to Pro-84, Asp-90
	to Ser-103, Gly-172 to Glu-179, Pro-190 to Phe-197, Val-210 to Arg-
	216, Pro-228 to Leu-233.
900777	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6370 as residues: Pro-5 to Arg-16, Thr-21 to Gly-27, Ser-35 to
	Gln-40, Arg-103 to Lys-112, Gly-172 to Pro-188, Gln-190 to Met-198.
900784	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6371 as residues: Gln-36 to Trp-52, Gly-164 to Gly-175, Ile-
	210 to Arg-215, Asn-417 to Val-422, Val-426 to Gln-431, Val-439 to
	Gly-444, Lys-470 to Leu-481, Phe-500 to Ser-511, Met-553 to Gly-563,
	Glu-691 to Thr-700, Ile-714 to Gly-723, Ala-750 to Gly-762, Leu-788 to

	
	Phe-794, Ser-798 to Gln-803, Thr-811 to Lys-816, Ser-824 to Phe-835, Thr-882 to Glu-892, Leu-901 to Gln-907, Gln-937 to Met-944.
900838	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6372 as residues: Pro-9 to Gly-15, Pro-47 to Pro-69, Pro-113 to Cys-122.
900966	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6374 as residues: Arg-34 to Gly-42, Gly-53 to Ser-59, Ala-74 to Gly-81, Glu-89 to Gly-103, Gly-108 to Gly-113, His-120 to Gly-223, Asp-225 to Gly-243, Pro-247 to Gly-312, Gly-317 to Asp-322.
901111	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6377 as residues: Pro-17 to Asp-36, Pro-102 to Glu-108, Pro-122 to Lys-128, His-150 to Gly-155, Asn-162 to Tyr-168, Pro-186 to Gln-193, Ser-205 to Pro-211, Gln-305 to Gly-317.
901128	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6379 as residues: Pro-1 to Gly-8, Pro-38 to Pro-45, Thr-103 to Ser-109, Cys-112 to Trp-119, Ala-201 to His-210, Glu-230 to Asn-241, Trp-263 to Ala-269.
901202	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6380 as residues: Pro-1 to Leu-17, Gly-36 to Gly-49.
901253	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6381 as residues: Gly-13 to Met-26, Arg-34 to Gly-39, Ile-60 to Ser-80, Ala-85 to Thr-98.
901276	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6382 as residues: Gln-1 to Arg-24, Gln-41 to Ala-48, Ser-70 to Gly-82, Glu-104 to Phe-112, Lys-126 to Ser-132, Pro-276 to Ile-281.
901333	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6383 as residues: Gln-48 to Lys-64, Glu-175 to Thr-183.
901375	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6384 as residues: Pro-3 to Lys-8, Phe-43 to Gly-51, Lys-55 to Ala-62, Ser-92 to Gln-98, Asp-106 to Trp-113, Ser-125 to Asn-134, Ser-150 to Phe-160.
901421	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6386 as residues: Arg-29 to Leu-38, Lys-47 to Arg-53, Asp-70 to Thr-75, Glu-116 to Leu-124, Gln-134 to Ser-143, Ser-158 to Trp-163, Pro-168 to Asp-180.
901472	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6387 as residues: Arg-1 to Val-7, Ala-156 to Phe-162.
901473	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6388 as residues: Leu-39 to Ile-47, Val-92 to Arg-98, Tyr-146 to Leu-160, Asp-185 to Phe-192, Phe-195 to Gly-207.
901494	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6389 as residues: Pro-11 to Trp-16, Gln-25 to Ser-37, Pro-99 to Gly-104, Pro-109 to Gly-115, Trp-201 to Thr-209.
901515	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6390 as residues: Gln-46 to Leu-51, Asp-58 to Asn-65, Lys-70 to Gln-75, Pro-111 to Thr-117, Gly-176 to Gly-185, Asp-205 to Gly-213, Thr-247 to Ile-263, Leu-269 to Lys-279.
901567	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6391 as residues: Phe-3 to Ala-8, Pro-17 to Gly-24, Asn-162 to Gln-179, Asn-195 to Asp-201, Glu-207 to Leu-213.

901578	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6392 as residues: Leu-1 to Glu-13, lle-34 to Arg-40, Lys-46 to
	Arg-57, Ala-77 to Ile-88, Pro-103 to Asp-111, Phe-127 to Ser-138.
901621	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6393 as residues: Gln-7 to Gly-12, Leu-60 to Pro-65, Arg-85 to
	Lys-99, Ser-132 to Pro-145, Pro-150 to Asp-155, Pro-183 to Asn-193,
	Arg-200 to Tyr-206.
901875	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6394 as residues: Gly-13 to Met-26, Arg-34 to Gly-39, Ile-60 to
	Ser-80, Ala-85 to Thr-98, Asn-109 to Val-140, Lys-150 to Thr-157, Gly-
	174 to Ala-201, Thr-204 to Lys-212, Thr-237 to Gly-243, Pro-251 to
	Pro-261, Ala-263 to Lys-277, Phe-281 to Arg-286, Arg-333 to Asp-341,
3	Glu-407 to Asp-412, Gly-424 to Gly-430, Gly-570 to Trp-583, Gln-614
	to Gly-619.
HCRMU56	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6396 as residues: Leu-7 to Leu-13, Pro-15 to Gln-27.
HKCSA70R	Preferred epitopes include those comprising a sequence shown in SEQ
THUI OD 10	ID NO. 6398 as residues: Leu-29 to Val-34, Gln-42 to Gly-52.
HWLOB10	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6399 as residues: Gly-49 to Pro-54.
HCQCG26R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6400 as residues: Gly-1 to Asp-6, Asp-16 to Ser-21, Val-36 to
HOENF69R	Cys-43, Ser-51 to Leu-60, Ile-65 to Lys-70.
HOENFOSK	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6404 as residues: Ala-15 to Ser-32, Ser-34 to Gly-43, Thr-57 to Gly-65.
HWLQY33	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6406 as residues: Gln-17 to Lys-22.
HCRNF08R	Preferred epitopes include those comprising a sequence shown in SEQ
Inola in out	ID NO. 6407 as residues: Arg-1 to Arg-13, Asn-33 to Arg-39.
HKCSZ69R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6408 as residues: Thr-32 to Lys-37.
HCQAG23	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6409 as residues: Arg-22 to Thr-28.
H2LAF75R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6411 as residues: Gly-1 to Ser-6.
H2LAT73R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6413 as residues: Thr-3 to Ser-10.
HUUAQ45	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6422 as residues: Arg-13 to Asn-22, Lys-42 to Glu-48.
HWLWQ51	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6423 as residues: Ala-18 to Asn-24, Thr-65 to Arg-71, Val-84
	to Thr-96.
HKLAB44R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6424 as residues: Val-7 to Trp-19, Ser-73 to Ser-79, Lys-86 to
11005:005	Ser-94.
H2CBA06R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6425 as residues: Ala-12 to Asp-20, Glu-30 to Arg-40, Gln-51 to Arg-57, Arg-79 to Tyr-88.

HCNAH60 R	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6427 as residues: Arg-19 to Gly-32.
HWMBJ68 R	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6432 as residues: Glu-10 to Gly-16, Asp-62 to Arg-69.
HELGR96R	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6435 as residues: Leu-31 to Gln-39.
HCRQM72 R	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6437 as residues: Asn-5 to Lys-14, Glu-25 to Gly-33, Arg-48 to Thr-74.
HWLMH52 R	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6440 as residues: Glu-24 to Leu-30.
H2CBU03R	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6441 as residues: Thr-2 to Ser-9.
HCQDR91R	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6443 as residues: Gly-14 to Arg-19.
HWMBN34 R	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6444 as residues: Lys-7 to Thr-12, Pro-25 to Lys-30, Leu-38 to Asp-43, Ser-84 to Ala-95, Asp-108 to Ser-117.
HCRNF81R	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6447 as residues: Pro-12 to His-17, Gln-57 to Asp-62, Thr-79 to Lys-101, Thr-117 to Ser-129.
HOHCI31R	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6448 as residues: Leu-16 to Ser-22, Lys-24 to Glu-38.
HSKKC10R	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6449 as residues: Glu-4 to Gly-10.
H2CBC52R	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6452 as residues: Pro-18 to Ser-30, Pro-37 to Pro-43.
HWLMC24 R	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6454 as residues: Pro-4 to Gly-34.
HWLUR40 R	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6455 as residues: Phe-3 to Lys-12.
HHAOD46 R	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6457 as residues: Lys-23 to Ala-40, Pro-67 to Ala-72, Val-102 to Thr-110.
HCYBA83R	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6458 as residues: Trp-13 to Ile-21, Pro-59 to Thr-68, Ala-85 to Lys-92, Thr-102 to Gly-113.
HCROZ77R	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6459 as residues: Asp-1 to Arg-8, Lys-13 to Leu-18, Gly-32 to Glu-49, Lys-60 to Ala-75, Ser-84 to Asp-99, Glu-107 to Ser-119, Ala-132 to Gly-141.
HCQCP20R	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6461 as residues: Leu-18 to Gln-25, Lys-37 to Phe-45.
HWLNF84 R	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6463 as residues: Lys-17 to Asn-22, Glu-31 to Lys-36, Gln-38 to Arg-44, Thr-81 to Thr-88.
HCRQI10R	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6468 as residues: Asp-56 to Lys-63, Lys-78 to Asn-86, Phe-92 to Lys-99.

HULCD94R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6470 as residues: Lys-7 to Thr-13, Asp-24 to Thr-30, Gly-39 to
	Glu-52, Leu-70 to Arg-76, Phe-87 to Tyr-92.
HHMMF84	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6471 as residues: Lys-30 to His-37.
HCRPO08R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6472 as residues: Val-33 to Lys-38.
HWLMQ74	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6475 as residues: Pro-9 to Gly-21.
H2LAB80R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6478 as residues: Thr-14 to Val-32.
HCQDO33	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6480 as residues: Trp-10 to Gly-18, Arg-34 to Pro-39.
HKAFL06R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6482 as residues: Pro-1 to Gly-14, Cys-18 to Gly-24, Ala-39 to
	Arg-55, Gly-63 to Glu-76, Gln-106 to Arg-115.
HWLOO35	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6486 as residues: Gly-1 to Gly-7, Arg-13 to Glu-19.
HWLVL77	Preferred epitopes include those comprising a sequence shown in SEQ
R HBJMG15R	ID NO. 6487 as residues: Arg-13 to Gly-40.
HRIMGISK	Preferred epitopes include those comprising a sequence shown in SEQ
H2CBH29R	ID NO. 6489 as residues: Ser-14 to Glu-27, Ile-40 to Ile-54.
H2CBH29K	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6490 as residues: Ser-16 to Glu-21.
H2LBB21R	
HZLBBZIK	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6494 as residues: Phe-50 to Tyr-55, Thr-63 to Trp-69, Pro-74 to
	Arg-80.
H2LAT69R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6495 as residues: Thr-2 to Ser-11.
HLWCJ40R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6496 as residues: Tyr-28 to Pro-40.
HOGDQ57	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6498 as residues: Pro-1 to Gln-8, Met-20 to Leu-26, Gly-42 to
	Ser-49, Ile-63 to Pro-73, Gly-80 to Ala-87.
HWLQM12	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6499 as residues: Pro-45 to Gly-52.
H2CBG89R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6501 as residues: Met-2 to Asp-31, Leu-67 to Asp-74, Gly-93 to
1	Ser-98.
HWLWQ68	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6502 as residues: Ser-21 to Glu-38.
НСҮВМ79	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6503 as residues: Glu-11 to Lys-22, Asp-31 to Trp-50.
HMUBO53	Preferred epitopes include those comprising a sequence shown in SEQ
RA	ID NO. 6504 as residues: Glu-1 to Asp-6, Asn-92 to Leu-97.
HWLVN81	Preferred epitopes include those comprising a sequence shown in SEQ
R HWI BV71	ID NO. 6506 as residues: Arg-6 to Val-14.
HWLRV71	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6507 as residues: Asp-34 to Pro-45.

Arg-73, Lys-87 to Phe-95. HWLNJ72R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6512 as residues: Ala-21 to Pro-30, Thr-43 to Glu-51. HOFME52 R BONO. 6513 as residues: Pro-7 to Phe-14, Glu-22 to Lys-28, Ala-31 to Glu-39, Lys-47 to Asp-54. HCRMG55 R BONO. 6515 as residues: Pro-7 to Phe-14, Glu-22 to Lys-28, Ala-31 to Glu-39, Lys-47 to Asp-54. HCRMG55 R BONO. 6515 as residues: Pro-4 to Gly-10, Lys-28 to Thr-37, Glu-45 to Glu-55. HCRNZ49R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6516 as residues: Pro-1 to Ala-14. H2LAD43R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6516 as residues: Gly-1 to Ser-6, Pro-20 to Trp-31. HCQCB53R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6522 as residues: Pro-8 to Asn-18. HCQCP47R HCQCP47R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6525 as residues: Thr-4 to Ser-11. HCQDC76R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6526 as residues: Thr-4 to Ser-11. HCQDC76R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6526 as residues: Gly-1 to Gly-8. HCQDK53 R HCQDK53 R HCQDK53 R HCQDK53 R HCQDK53 R HCQDK54 R HCQDK54 R HCQDK55 R HCQDK55 R HCQDK55 R HCQDK55 R HCQDK50 R HCQDK5	HDPMJ48R	Preferred epitopes include those comprising a sequence shown in SEQ
HWLNJ72R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6512 as residues: Ala-21 to Pro-30, Thr-43 to Glu-51. HOFME52 R Dreferred epitopes include those comprising a sequence shown in SEQ ID NO. 6513 as residues: Pro-7 to Phe-14, Glu-22 to Lys-28, Ala-31 to Glu-39, Lys-47 to Asp-54. HCRMG55 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6515 as residues: Pro-4 to Gly-10, Lys-28 to Thr-37, Glu-45 to Glu-55. HCRNZ49R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6516 as residues: Pro-1 to Ala-14. H2LAD43R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6518 as residues: Gly-1 to Ser-6, Pro-20 to Trp-31. HCQCB53R HCQCL32R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6522 as residues: Pro-8 to Asn-18. HCQCL32R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6525 as residues: Arg-3 to Asn-18. HCQCP47R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6526 as residues: Thr-4 to Ser-11. HCQDC76R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6526 as residues: Thr-4 to Ser-11. HCQDC76R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6528 as residues: Gly-1 to Lys-6, Lys-11 to Ser-17. HCQDK53 R HCQDK53 R HCQDK53 R HCQDK53 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6538 as residues: Gly-1 to Gly-8. HCQDK54 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6533 as residues: Gly-1 to Gly-8. HCQDK55 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6543 as residues: Gly-1 to Gly-8. HCQDF48 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6553 as residues: Ser-31 to Tyr-36, Pro-64 to Gly-72. HCRNF48R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6555 as residues: Ser-31 to Tyr-36, Pro-64 to Gly-72. HCRDF34R Preferred epitopes inclu		ID NO. 6511 as residues: Thr-1 to Trp-14, Lys-27 to Leu-44, Glu-59 to
HOFMES2 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6516 as residues: Pro-7 to Phe-14, Glu-22 to Lys-28, Ala-31 to Glu-39, Lys-47 to Asp-54. HCRMG55 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6516 as residues: Pro-4 to Gly-10, Lys-28 to Thr-37, Glu-45 to Glu-55. HCRNZ49R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6516 as residues: Pro-1 to Ala-14. H2LAD43R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6518 as residues: Gly-1 to Ser-6, Pro-20 to Trp-31. HCQCB53R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6522 as residues: Pro-8 to Asn-18. HCQCL32R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6525 as residues: Arg-3 to Asn-18. HCQCP47R HCQCP47R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6526 as residues: Arg-3 to Asn-18. HCQCP47R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6526 as residues: Arg-1 to Lys-6, Lys-11 to Ser-17. HCQDT68 R HCQDK53 R HCQDK53 R HCQDK53 R HCQDK53 R HCQDF62 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6528 as residues: Gly-1 to Gly-8. HCQDF62 R HCQDF63 R HCQDF63 R HCQDF63 R HCQDF64 R DNO. 6535 as residues: Gly-1 to Gly-8. HCQDF654 R HCQDF655 R HCQDF67 R HCQDF68 R HCQDF68 R HCQDF68 R HCQDF68 R HCQDF69 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6535 as residues: Gly-1 to Gly-8. HCQDF69 R HCRNF458 R HCQDF69 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6536 as residues: Ser-31 to Tyr-36, Pro-64 to Gly-72. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6556 as residues: Ser-63 to Gly-69. HCRNF458 R HCRNF458 R HCRNF458 R HCRNF458 R HCRNF458 R HCRNF458 R HCRNF458 R HCRNF458 R HCRNF458 R HCRNF458 R HCRNF458 R HCRNF458 R HCRNF468 R HCRNF468 R HCRNF47 R HCRNF47 R HCRNF47 R HCRNF47 R HCRNF47 R HCRNF47 R HCRNF47 R HC	111111 > 1120	
HOFME52 R DNO. 6513 as residues: Pro-7 to Phe-14, Glu-22 to Lys-28, Ala-31 to Glu-39, Lys-47 to Asp-54. HCRMG55 R DNO. 6515 as residues: Pro-7 to Phe-14, Glu-22 to Lys-28, Ala-31 to Glu-39, Lys-47 to Asp-54. HCRMG55 R DNO. 6515 as residues: Pro-4 to Gly-10, Lys-28 to Thr-37, Glu-45 to Glu-55. HCRNZ49R Dreferred epitopes include those comprising a sequence shown in SEQ ID NO. 6518 as residues: Pro-1 to Ala-14. H2LAD43R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6518 as residues: Gly-1 to Ser-6, Pro-20 to Trp-31. HCQCB53R HCQCB53R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6522 as residues: Pro-8 to Asn-18. HCQCL32R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6525 as residues: Arg-3 to Asn-18. HCQCP47R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6526 as residues: Thr-4 to Ser-11. HCQDC76R HCQDC76R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6527 as residues: Thr-4 to Ser-11. HCQDK53 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6528 as residues: Gly-1 to Gly-8. HCQDK53 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6531 as residues: Gly-1 to Gly-8. HCQDF6R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6533 as residues: Gly-1 to Gly-8. HCQDF6R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6533 as residues: Gly-1 to Gly-8. HCQDF6R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6533 as residues: Gly-1 to Gly-8. HCQDF6R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6535 as residues: Pro-8 to Glu-13, Pro-27 to Pro-33. HCRNF45R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6556 as residues: Pro-8 to Glu-13, Pro-27 to Pro-33. HCRNF45R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6556 as residues: Ser-4 to Pro-10, Glu-18 to	HWLNJ/2R	
R ID NO. 6513 as residues: Pro-7 to Phe-14, Glu-22 to Lys-28, Ala-31 to Glu-39, Lys-47 to Asp-54. HCRMG55 R Dreferred epitopes include those comprising a sequence shown in SEQ ID NO. 6516 as residues: Pro-4 to Gly-10, Lys-28 to Thr-37, Glu-45 to Glu-55. HCRNZ49R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6516 as residues: Pro-1 to Ala-14. H2LAD43R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6518 as residues: Gly-1 to Ser-6, Pro-20 to Trp-31. HCQCB53R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6522 as residues: Pro-8 to Asn-18. HCQCL32R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6525 as residues: Arg-3 to Asn-18. HCQCP47R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6526 as residues: Thr-4 to Ser-11. HCQDC76R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6527 as residues: Asp-1 to Lys-6, Lys-11 to Ser-17. HCQDH59 R DNO. 6528 as residues: Gly-1 to Lys-6, Lys-11 to Ser-17. HCQDK53 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6528 as residues: Gly-1 to Gly-8. HCQDF62R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6531 as residues: Gly-1 to Gly-8. HCQDF62R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6533 as residues: Gly-1 to Gly-8. HCQDF63R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6534 as residues: Ser-31 to Tyr-36, Pro-64 to Gly-72. HCRNF45R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6546 as residues: Pro-8 to Glu-13, Pro-27 to Pro-33. HCROB90R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6556 as residues: Arg-63 to Gly-69. HCRNF45R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6556 as residues: Ser-15 to Ile-24, Asn-56 to Lys-67, Ser-80 to Lys-95. HCRDF3R Preferred epitopes include those comprising a s	HOD (DIO	
Glu-39, Lys-47 to Asp-54. HCRMG55 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6515 as residues: Pro-4 to Gly-10, Lys-28 to Thr-37, Glu-45 to Glu-55. HCRNZ49R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6516 as residues: Pro-1 to Ala-14. H2LAD43R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6518 as residues: Gly-1 to Ser-6, Pro-20 to Trp-31. HCQCB53R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6522 as residues: Pro-8 to Asn-18. HCQCL32R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6525 as residues: Arg-3 to Asn-18. HCQCP47R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6526 as residues: Thr-4 to Ser-11. HCQDC76R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6527 as residues: Asp-1 to Lys-6, Lys-11 to Ser-17. HCQDH59 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6528 as residues: Gly-1 to Gly-8. HCQDK53 R HCQDK53 R HCQDF62R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6531 as residues: Gly-1 to Gly-8. HCQDP62R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6533 as residues: Gly-1 to Gly-8. HCQDF63R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6543 as residues: Ser-31 to Tyr-36, Pro-64 to Gly-72. HCRNF45R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6546 as residues: Pro-8 to Glu-13, Pro-27 to Pro-33. HCROB90R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6556 as residues: Pro-8 to Glu-13, Pro-27 to Pro-33. HCRNIS0R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6557 as residues: Ser-31 to Tyr-36, Pro-64 to Gly-72. HCRPJ34R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6555 as residues: Ser-4 to Pro-10, Glu-18 to Cys-23. HVLOR95 Preferred epitopes include tho	·	
HCRMG55 R DNO. 6515 as residues: Pro-4 to Gly-10, Lys-28 to Thr-37, Glu-45 to Glu-55. HCRNZ49R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6516 as residues: Pro-1 to Ala-14. H2LAD43R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6518 as residues: Gly-1 to Ser-6, Pro-20 to Trp-31. HCQCB53R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6522 as residues: Pro-8 to Asn-18. HCQCL32R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6525 as residues: Arg-3 to Asn-18. HCQCP47R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6525 as residues: Arg-3 to Asn-18. HCQDC76R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6527 as residues: Asp-1 to Lys-6, Lys-11 to Ser-17. HCQDC76R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6527 as residues: Gly-1 to Gly-8. HCQDK53 R HCQDK53 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6531 as residues: Gly-1 to Gly-8. HCQDP62R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6535 as residues: Gly-1 to Gly-8. HKCAA76 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6535 as residues: Gly-1 to Gly-8. HCRNF45R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6546 as residues: Ser-31 to Tyr-36, Pro-64 to Gly-72. HCRNF45R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6556 as residues: Ser-31 to Tyr-36, Pro-64 to Lys-67, Ser-80 to Lys-95. HCRNI50R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6555 as residues: Ser-15 to Ile-24, Asn-56 to Lys-67, Ser-80 to Lys-95. HCRDJ34R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6556 as residues: Ser-15 to Ile-24, Asn-56 to Lys-67, Ser-80 to Lys-95. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6556 as residues: Ser-4 to Pro-10, G	R	
R ID NO. 6515 as residues: Pro-4 to Gly-10, Lys-28 to Thr-37, Glu-45 to Glu-55. HCRNZ49R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6516 as residues: Pro-1 to Ala-14. H2LAD43R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6518 as residues: Gly-1 to Ser-6, Pro-20 to Trp-31. HCQCB53R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6522 as residues: Pro-8 to Asn-18. HCQCL32R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6525 as residues: Arg-3 to Asn-18. HCQCP47R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6526 as residues: Thr-4 to Ser-11. HCQDC76R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6527 as residues: Asp-1 to Lys-6, Lys-11 to Ser-17. HCQDH59 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6528 as residues: Gly-1 to Gly-8. HCQDK53 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6531 as residues: Gly-1 to Gly-8. HCQDP62R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6535 as residues: Gly-1 to Gly-8. HKCAA76 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6543 as residues: Ser-31 to Tyr-36, Pro-64 to Gly-72. HCRNF45R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6546 as residues: Pro-8 to Glu-13, Pro-27 to Pro-33. HCROB90R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6555 as residues: Arg-63 to Gly-69. HCRNI50R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6556 as residues: Ser-15 to Ile-24, Asn-56 to Lys-67, Ser-80 to Lys-95. HCRDJ34R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6555 as residues: Ser-15 to Ile-24, Asn-56 to Lys-67, Ser-80 to Lys-95. HCRDJ34R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6556 as residues: Ser-4 to Pro-10, Glu-18 to Cys-23. HCCQBL9	HCRMG55	
HCRNZ49R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6516 as residues: Pro-1 to Ala-14. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6518 as residues: Gly-1 to Ser-6, Pro-20 to Trp-31. HCQCB53R	R	ID NO. 6515 as residues: Pro-4 to Gly-10, Lys-28 to Thr-37, Glu-45 to
H2LAD43R H2LAD43R H7eferred epitopes include those comprising a sequence shown in SEQ ID NO. 6518 as residues: Gly-1 to Ser-6, Pro-20 to Trp-31. HCQCB53R HCQCL32R HCQCL32R HCQCL32R HCQCL32R HCQCP47R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6522 as residues: Pro-8 to Asn-18. HCQCP47R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6525 as residues: Arg-3 to Asn-18. HCQDC76R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6526 as residues: Thr-4 to Ser-11. HCQDH59 R HCQDH59 R HCQDK53 R HCQDK53 R HCQDK53 R HCQDF62R HCQDC64R HCQDC64R HCQDC65B HCQCC65B HCCC65B	TICDNIZ 40D	
D NO. 6518 as residues: Gly-1 to Ser-6, Pro-20 to Trp-31. HCQCB53R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6522 as residues: Pro-8 to Asn-18. HCQCL32R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6525 as residues: Arg-3 to Asn-18. HCQCP47R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6526 as residues: Thr-4 to Ser-11. HCQDC76R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6526 as residues: Asp-1 to Lys-6, Lys-11 to Ser-17. HCQDH59 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6528 as residues: Gly-1 to Gly-8. HCQDK53 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6531 as residues: Gly-1 to Gly-8. HCQDP62R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6535 as residues: Gly-1 to Gly-8. HKCAA76 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6543 as residues: Ser-31 to Tyr-36, Pro-64 to Gly-72. HCRNF45R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6543 as residues: Pro-8 to Glu-13, Pro-27 to Pro-33. HCROB90R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6555 as residues: Arg-63 to Gly-69. HCRNI50R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6556 as residues: Ser-15 to Ile-24, Asn-56 to Lys-67, Ser-80 to Lys-95. HCRPJ34R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6556 as residues: Ser-15 to Ile-24, Asn-56 to Lys-67, Ser-80 to Lys-95. HCRPJ34R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6556 as residues: Ser-4 to Pro-10, Glu-18 to Cys-23. HWLOR95 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6559 as residues: Ser-23 to Ala-28, Pro-64 to Glu-74, Ala-100 to Lys-106. HKCS132R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6560 as residues: Ala-4 to Gln-14, Gly-3	HCKNZ49K	
HCQCB53R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6522 as residues: Pro-8 to Asn-18. HCQCL32R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6525 as residues: Arg-3 to Asn-18. HCQCP47R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6526 as residues: Thr-4 to Ser-11. HCQDC76R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6527 as residues: Asp-1 to Lys-6, Lys-11 to Ser-17. HCQDH59 R ID NO. 6528 as residues: Gly-1 to Gly-8. HCQDK53 R HCQDK53 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6531 as residues: Gly-1 to Gly-8. HCQDP62R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6533 as residues: Gly-1 to Gly-8. HKCAA76 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6543 as residues: Ser-31 to Tyr-36, Pro-64 to Gly-72. HCRNF45R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6546 as residues: Pro-8 to Glu-13, Pro-27 to Pro-33. HCROB90R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6555 as residues: Arg-63 to Gly-69. HCRNI50R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6556 as residues: Ser-15 to Ile-24, Asn-56 to Lys-67, Ser-80 to Lys-95. HCRPJ34R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6556 as residues: Ser-15 to Ile-24, Asn-56 to Lys-67, Ser-80 to Lys-95. HCRPJ34R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6558 as residues: Ser-4 to Pro-10, Glu-18 to Cys-23. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6559 as residues: Ser-23 to Ala-28, Pro-64 to Glu-74, Ala-100 to Lys-106. HKCS132R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6560 as residues: Ala-4 to Gln-14, Gly-36 to Gln-42, Gly-70 to Leu-77.	H2LAD43R	Preferred epitopes include those comprising a sequence shown in SEQ
ID NO. 6522 as residues: Pro-8 to Asn-18. HCQCL32R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6525 as residues: Arg-3 to Asn-18. HCQCP47R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6526 as residues: Thr-4 to Ser-11. HCQDC76R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6527 as residues: Asp-1 to Lys-6, Lys-11 to Ser-17. HCQDH59 R ID NO. 6528 as residues: Gly-1 to Gly-8. HCQDK53 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6531 as residues: Gly-1 to Gly-8. HCQDP62R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6535 as residues: Gly-1 to Gly-8. HKCAA76 R ID NO. 6535 as residues: Ser-31 to Tyr-36, Pro-64 to Gly-72. HCRNF45R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6546 as residues: Pro-8 to Glu-13, Pro-27 to Pro-33. HCROB90R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6555 as residues: Arg-63 to Gly-69. HCRNISOR HCRNISOR Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6556 as residues: Ser-15 to Ile-24, Asn-56 to Lys-67, Ser-80 to Lys-95. HCRPJ34R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6557 as residues: Ser-15 to Ile-24, Asn-56 to Lys-67, Ser-80 to Lys-95. HCRPJ34R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6557 as residues: Ser-4 to Pro-10, Glu-18 to Cys-23. HWLOR95 R ID NO. 6559 as residues: Ser-4 to Pro-10, Glu-18 to Cys-23. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6559 as residues: Ser-4 to Pro-64 to Glu-74, Ala-100 to Lys-106. HKCS132R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 65560 as residues: Ser-23 to Ala-28, Pro-64 to Glu-74, Ala-100 to Lys-106. HKCS132R	\	ID NO. 6518 as residues: Gly-1 to Ser-6, Pro-20 to Trp-31.
ID NO. 6522 as residues: Pro-8 to Asn-18. HCQCL32R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6525 as residues: Arg-3 to Asn-18. HCQCP47R HCQDC76R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6526 as residues: Thr-4 to Ser-11. HCQDC76R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6527 as residues: Asp-1 to Lys-6, Lys-11 to Ser-17. HCQDH59 R ID NO. 6528 as residues: Gly-1 to Gly-8. HCQDK53 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6531 as residues: Gly-1 to Gly-8. HCQDP62R HCQDP62R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6535 as residues: Gly-1 to Gly-8. HKCAA76 R ID NO. 6535 as residues: Ser-31 to Tyr-36, Pro-64 to Gly-72. HCRNF45R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6546 as residues: Pro-8 to Glu-13, Pro-27 to Pro-33. HCROB90R HCROB90R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6555 as residues: Arg-63 to Gly-69. HCRNI50R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6556 as residues: Ser-15 to Ile-24, Asn-56 to Lys-67, Ser-80 to Lys-95. HCRPJ34R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6557 as residues: Ser-15 to Ile-24, Asn-56 to Lys-67, Ser-80 to Lys-95. HCRPJ34R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6557 as residues: Ser-4 to Pro-10, Glu-18 to Cys-23. HWLOR95 R ID NO. 6559 as residues: Ser-4 to Pro-10, Glu-18 to Cys-23. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6559 as residues: Ser-4 to Pro-10, Glu-18 to Cys-23. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6559 as residues: Ser-23 to Ala-28, Pro-64 to Glu-74, Ala-100 to Lys-106. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6550 as residues: Ser-23 to Ala-28, Pro-64 to Glu-74, Ala-100 to Lys-106.	HCQCB53R	Preferred epitopes include those comprising a sequence shown in SEQ
HCQCP47R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6526 as residues: Thr-4 to Ser-11. HCQDC76R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6527 as residues: Asp-1 to Lys-6, Lys-11 to Ser-17. HCQDH59 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6528 as residues: Gly-1 to Gly-8. HCQDK53 R HCQDK53 R HCQDP62R D NO. 6531 as residues: Gly-1 to Gly-8. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6535 as residues: Gly-1 to Gly-8. HKCAA76 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6535 as residues: Gly-1 to Gly-8. HKCAA76 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6543 as residues: Ser-31 to Tyr-36, Pro-64 to Gly-72. HCRNF45R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6546 as residues: Pro-8 to Glu-13, Pro-27 to Pro-33. HCROB90R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6555 as residues: Arg-63 to Gly-69. HCRNI50R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6556 as residues: Ser-15 to Ile-24, Asn-56 to Lys-67, Ser-80 to Lys-95. HCRPJ34R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6557 as residues: Val-5 to Gln-11. HCQBL95R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6559 as residues: Ser-4 to Pro-10, Glu-18 to Cys-23. HWLOR95 R ID NO. 6559 as residues: Ser-23 to Ala-28, Pro-64 to Glu-74, Ala-100 to Lys-106. HKCSI32R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6550 as residues: Ser-23 to Ala-28, Pro-64 to Glu-74, Ala-100 to Lys-106. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6550 as residues: Ala-4 to Gln-14, Gly-36 to Gln-42, Gly-70 to Leu-77.		ID NO. 6522 as residues: Pro-8 to Asn-18.
HCQCP47R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6526 as residues: Thr-4 to Ser-11. HCQDC76R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6527 as residues: Asp-1 to Lys-6, Lys-11 to Ser-17. HCQDH59 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6528 as residues: Gly-1 to Gly-8. HCQDK53 R HCQDK53 R HCQDF62R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6531 as residues: Gly-1 to Gly-8. HCQDP62R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6535 as residues: Gly-1 to Gly-8. HKCAA76 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6543 as residues: Ser-31 to Tyr-36, Pro-64 to Gly-72. HCRNF45R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6546 as residues: Pro-8 to Glu-13, Pro-27 to Pro-33. HCROB90R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6555 as residues: Arg-63 to Gly-69. HCRNI50R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6556 as residues: Ser-15 to Ile-24, Asn-56 to Lys-67, Ser-80 to Lys-95. HCRPJ34R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6557 as residues: Val-5 to Gln-11. HCQBL95R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6559 as residues: Ser-4 to Pro-10, Glu-18 to Cys-23. HWLOR95 R ID NO. 6559 as residues: Ser-23 to Ala-28, Pro-64 to Glu-74, Ala-100 to Lys-106. HKCSI32R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6550 as residues: Ser-23 to Ala-28, Pro-64 to Glu-74, Ala-100 to Lys-106. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6550 as residues: Ala-4 to Gln-14, Gly-36 to Gln-42, Gly-70 to Leu-77.	HCQCL32R	Preferred epitopes include those comprising a sequence shown in SEQ
ID NO. 6526 as residues: Thr-4 to Ser-11. HCQDC76R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6527 as residues: Asp-1 to Lys-6, Lys-11 to Ser-17. HCQDH59 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6528 as residues: Gly-1 to Gly-8. HCQDK53 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6531 as residues: Gly-1 to Gly-8. HCQDP62R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6535 as residues: Gly-1 to Gly-8. HKCAA76 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6543 as residues: Ser-31 to Tyr-36, Pro-64 to Gly-72. HCRNF45R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6546 as residues: Pro-8 to Glu-13, Pro-27 to Pro-33. HCROB90R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6555 as residues: Arg-63 to Gly-69. HCRNI50R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6556 as residues: Ser-15 to Ile-24, Asn-56 to Lys-67, Ser-80 to Lys-95. HCRPJ34R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6557 as residues: Val-5 to Gln-11. HCQBL95R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6558 as residues: Ser-4 to Pro-10, Glu-18 to Cys-23. HWLOR95 R ID NO. 6559 as residues: Ser-23 to Ala-28, Pro-64 to Glu-74, Ala-100 to Lys-106. HKCSI32R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6560 as residues: Ala-4 to Gln-14, Gly-36 to Gln-42, Gly-70 to Leu-77.		ID NO. 6525 as residues: Arg-3 to Asn-18.
ID NO. 6526 as residues: Thr-4 to Ser-11. HCQDC76R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6527 as residues: Asp-1 to Lys-6, Lys-11 to Ser-17. HCQDH59 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6528 as residues: Gly-1 to Gly-8. HCQDK53 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6531 as residues: Gly-1 to Gly-8. HCQDP62R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6535 as residues: Gly-1 to Gly-8. HKCAA76 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6543 as residues: Ser-31 to Tyr-36, Pro-64 to Gly-72. HCRNF45R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6546 as residues: Pro-8 to Glu-13, Pro-27 to Pro-33. HCROB90R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6555 as residues: Arg-63 to Gly-69. HCRNI50R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6556 as residues: Ser-15 to Ile-24, Asn-56 to Lys-67, Ser-80 to Lys-95. HCRPJ34R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6557 as residues: Val-5 to Gln-11. HCQBL95R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6558 as residues: Ser-4 to Pro-10, Glu-18 to Cys-23. HWLOR95 R ID NO. 6559 as residues: Ser-23 to Ala-28, Pro-64 to Glu-74, Ala-100 to Lys-106. HKCSI32R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6560 as residues: Ala-4 to Gln-14, Gly-36 to Gln-42, Gly-70 to Leu-77.	HCQCP47R	Preferred epitopes include those comprising a sequence shown in SEQ
HCQDC76R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6527 as residues: Asp-1 to Lys-6, Lys-11 to Ser-17. HCQDH59 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6528 as residues: Gly-1 to Gly-8. HCQDK53 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6531 as residues: Gly-1 to Gly-8. HCQDP62R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6535 as residues: Gly-1 to Gly-8. HKCAA76 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6543 as residues: Ser-31 to Tyr-36, Pro-64 to Gly-72. HCRNF45R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6546 as residues: Pro-8 to Glu-13, Pro-27 to Pro-33. HCROB90R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6555 as residues: Arg-63 to Gly-69. HCRNI50R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6556 as residues: Ser-15 to Ile-24, Asn-56 to Lys-67, Ser-80 to Lys-95. HCRPJ34R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6557 as residues: Val-5 to Gln-11. HCQBL95R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6558 as residues: Ser-4 to Pro-10, Glu-18 to Cys-23. HWLOR95 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6559 as residues: Ser-23 to Ala-28, Pro-64 to Glu-74, Ala-100 to Lys-106. HKCSI32R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6560 as residues: Ala-4 to Gln-14, Gly-36 to Gln-42, Gly-70 to Leu-77.		ID NO. 6526 as residues: Thr-4 to Ser-11.
ID NO. 6527 as residues: Asp-1 to Lys-6, Lys-11 to Ser-17. HCQDH59 R ID NO. 6528 as residues: Gly-1 to Gly-8. HCQDK53 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6531 as residues: Gly-1 to Gly-8. HCQDP62R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6531 as residues: Gly-1 to Gly-8. HKCAA76 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6535 as residues: Gly-1 to Gly-8. HKCAA76 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6543 as residues: Ser-31 to Tyr-36, Pro-64 to Gly-72. HCRNF45R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6546 as residues: Pro-8 to Glu-13, Pro-27 to Pro-33. HCROB90R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6555 as residues: Arg-63 to Gly-69. HCRNI50R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6556 as residues: Ser-15 to Ile-24, Asn-56 to Lys-67, Ser-80 to Lys-95. HCRPJ34R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6557 as residues: Val-5 to Gln-11. HCQBL95R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6558 as residues: Ser-4 to Pro-10, Glu-18 to Cys-23. HWLOR95 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6559 as residues: Ser-23 to Ala-28, Pro-64 to Glu-74, Ala-100 to Lys-106. HKCSI32R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6560 as residues: Ala-4 to Gln-14, Gly-36 to Gln-42, Gly-70 to Leu-77.	HCQDC76R	
R ID NO. 6528 as residues: Gly-1 to Gly-8. HCQDK53 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6531 as residues: Gly-1 to Gly-8. HCQDP62R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6535 as residues: Gly-1 to Gly-8. HKCAA76 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6543 as residues: Ser-31 to Tyr-36, Pro-64 to Gly-72. HCRNF45R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6546 as residues: Pro-8 to Glu-13, Pro-27 to Pro-33. HCROB90R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6555 as residues: Arg-63 to Gly-69. HCRNI50R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6556 as residues: Ser-15 to Ile-24, Asn-56 to Lys-67, Ser-80 to Lys-95. HCRPJ34R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6557 as residues: Val-5 to Gln-11. HCQBL95R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6558 as residues: Ser-4 to Pro-10, Glu-18 to Cys-23. HWLOR95 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6559 as residues: Ser-23 to Ala-28, Pro-64 to Glu-74, Ala-100 to Lys-106. HKCS132R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6560 as residues: Ala-4 to Gln-14, Gly-36 to Gln-42, Gly-70 to Leu-77.		ID NO. 6527 as residues: Asp-1 to Lys-6, Lys-11 to Ser-17.
R ID NO. 6528 as residues: Gly-1 to Gly-8. HCQDK53 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6531 as residues: Gly-1 to Gly-8. HCQDP62R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6535 as residues: Gly-1 to Gly-8. HKCAA76 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6543 as residues: Ser-31 to Tyr-36, Pro-64 to Gly-72. HCRNF45R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6546 as residues: Pro-8 to Glu-13, Pro-27 to Pro-33. HCROB90R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6555 as residues: Arg-63 to Gly-69. HCRNI50R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6556 as residues: Ser-15 to Ile-24, Asn-56 to Lys-67, Ser-80 to Lys-95. HCRPJ34R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6557 as residues: Val-5 to Gln-11. HCQBL95R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6558 as residues: Ser-4 to Pro-10, Glu-18 to Cys-23. HWLOR95 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6559 as residues: Ser-23 to Ala-28, Pro-64 to Glu-74, Ala-100 to Lys-106. HKCS132R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6560 as residues: Ala-4 to Gln-14, Gly-36 to Gln-42, Gly-70 to Leu-77.	HCQDH59	
HCQDP62R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6535 as residues: Gly-1 to Gly-8. HKCAA76 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6543 as residues: Ser-31 to Tyr-36, Pro-64 to Gly-72. HCRNF45R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6546 as residues: Pro-8 to Glu-13, Pro-27 to Pro-33. HCROB90R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6555 as residues: Arg-63 to Gly-69. HCRNI50R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6556 as residues: Ser-15 to Ile-24, Asn-56 to Lys-67, Ser-80 to Lys-95. HCRPJ34R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6557 as residues: Val-5 to Gln-11. HCQBL95R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6558 as residues: Ser-4 to Pro-10, Glu-18 to Cys-23. HWLOR95 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6559 as residues: Ser-23 to Ala-28, Pro-64 to Glu-74, Ala-100 to Lys-106. HKCS132R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6560 as residues: Ala-4 to Gln-14, Gly-36 to Gln-42, Gly-70 to Leu-77.	, -	ID NO. 6528 as residues: Gly-1 to Gly-8.
HCQDP62R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6535 as residues: Gly-1 to Gly-8. HKCAA76 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6543 as residues: Ser-31 to Tyr-36, Pro-64 to Gly-72. HCRNF45R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6546 as residues: Pro-8 to Glu-13, Pro-27 to Pro-33. HCROB90R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6555 as residues: Arg-63 to Gly-69. HCRNI50R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6556 as residues: Ser-15 to Ile-24, Asn-56 to Lys-67, Ser-80 to Lys-95. HCRPJ34R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6557 as residues: Val-5 to Gln-11. HCQBL95R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6558 as residues: Ser-4 to Pro-10, Glu-18 to Cys-23. HWLOR95 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6559 as residues: Ser-23 to Ala-28, Pro-64 to Glu-74, Ala-100 to Lys-106. HKCSI32R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6560 as residues: Ala-4 to Gln-14, Gly-36 to Gln-42, Gly-70 to Leu-77.	HCQDK53	Preferred epitopes include those comprising a sequence shown in SEQ
HKCAA76 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6543 as residues: Ser-31 to Tyr-36, Pro-64 to Gly-72. HCRNF45R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6546 as residues: Pro-8 to Glu-13, Pro-27 to Pro-33. HCROB90R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6555 as residues: Arg-63 to Gly-69. HCRNI50R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6556 as residues: Ser-15 to Ile-24, Asn-56 to Lys-67, Ser-80 to Lys-95. HCRPJ34R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6557 as residues: Val-5 to Gln-11. HCQBL95R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6558 as residues: Ser-4 to Pro-10, Glu-18 to Cys-23. HWLOR95 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6559 as residues: Ser-23 to Ala-28, Pro-64 to Glu-74, Ala-100 to Lys-106. HKCSI32R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6560 as residues: Ala-4 to Gln-14, Gly-36 to Gln-42, Gly-70 to Leu-77.	, R	ID NO. 6531 as residues: Gly-1 to Gly-8.
HKCAA76 R Dreferred epitopes include those comprising a sequence shown in SEQ ID NO. 6543 as residues: Ser-31 to Tyr-36, Pro-64 to Gly-72. HCRNF45R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6546 as residues: Pro-8 to Glu-13, Pro-27 to Pro-33. HCROB90R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6555 as residues: Arg-63 to Gly-69. HCRNI50R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6556 as residues: Ser-15 to Ile-24, Asn-56 to Lys-67, Ser-80 to Lys-95. HCRPJ34R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6557 as residues: Val-5 to Gln-11. HCQBL95R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6558 as residues: Ser-4 to Pro-10, Glu-18 to Cys-23. HWLOR95 R ID NO. 6559 as residues: Ser-23 to Ala-28, Pro-64 to Glu-74, Ala-100 to Lys-106. HKCSI32R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6560 as residues: Ala-4 to Gln-14, Gly-36 to Gln-42, Gly-70 to Leu-77.	HCQDP62R	Preferred epitopes include those comprising a sequence shown in SEQ
HCRNF45R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6546 as residues: Pro-8 to Glu-13, Pro-27 to Pro-33. HCROB90R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6555 as residues: Arg-63 to Gly-69. HCRNI50R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6556 as residues: Ser-15 to Ile-24, Asn-56 to Lys-67, Ser-80 to Lys-95. HCRPJ34R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6557 as residues: Val-5 to Gln-11. HCQBL95R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6558 as residues: Ser-4 to Pro-10, Glu-18 to Cys-23. HWLOR95 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6559 as residues: Ser-23 to Ala-28, Pro-64 to Glu-74, Ala-100 to Lys-106. HKCSI32R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6560 as residues: Ala-4 to Gln-14, Gly-36 to Gln-42, Gly-70 to Leu-77.		ID NO. 6535 as residues: Gly-1 to Gly-8.
HCRNF45R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6546 as residues: Pro-8 to Glu-13, Pro-27 to Pro-33. HCROB90R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6555 as residues: Arg-63 to Gly-69. HCRNI50R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6556 as residues: Ser-15 to Ile-24, Asn-56 to Lys-67, Ser-80 to Lys-95. HCRPJ34R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6557 as residues: Val-5 to Gln-11. HCQBL95R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6558 as residues: Ser-4 to Pro-10, Glu-18 to Cys-23. HWLOR95 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6559 as residues: Ser-23 to Ala-28, Pro-64 to Glu-74, Ala-100 to Lys-106. HKCSI32R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6560 as residues: Ala-4 to Gln-14, Gly-36 to Gln-42, Gly-70 to Leu-77.	HKCAA76	Preferred epitopes include those comprising a sequence shown in SEQ
ID NO. 6546 as residues: Pro-8 to Glu-13, Pro-27 to Pro-33. HCROB90R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6555 as residues: Arg-63 to Gly-69. HCRNI50R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6556 as residues: Ser-15 to Ile-24, Asn-56 to Lys-67, Ser-80 to Lys-95. HCRPJ34R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6557 as residues: Val-5 to Gln-11. HCQBL95R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6558 as residues: Ser-4 to Pro-10, Glu-18 to Cys-23. HWLOR95 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6559 as residues: Ser-23 to Ala-28, Pro-64 to Glu-74, Ala-100 to Lys-106. HKCSI32R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6560 as residues: Ala-4 to Gln-14, Gly-36 to Gln-42, Gly-70 to Leu-77.	R	ID NO. 6543 as residues: Ser-31 to Tyr-36, Pro-64 to Gly-72.
HCROB90R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6555 as residues: Arg-63 to Gly-69. HCRNI50R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6556 as residues: Ser-15 to Ile-24, Asn-56 to Lys-67, Ser-80 to Lys-95. HCRPJ34R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6557 as residues: Val-5 to Gln-11. HCQBL95R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6558 as residues: Ser-4 to Pro-10, Glu-18 to Cys-23. HWLOR95 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6559 as residues: Ser-23 to Ala-28, Pro-64 to Glu-74, Ala-100 to Lys-106. HKCSI32R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6560 as residues: Ala-4 to Gln-14, Gly-36 to Gln-42, Gly-70 to Leu-77.	HCRNF45R	l
ID NO. 6555 as residues: Arg-63 to Gly-69. HCRNI50R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6556 as residues: Ser-15 to Ile-24, Asn-56 to Lys-67, Ser-80 to Lys-95. HCRPJ34R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6557 as residues: Val-5 to Gln-11. HCQBL95R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6558 as residues: Ser-4 to Pro-10, Glu-18 to Cys-23. HWLOR95 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6559 as residues: Ser-23 to Ala-28, Pro-64 to Glu-74, Ala-100 to Lys-106. HKCSI32R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6560 as residues: Ala-4 to Gln-14, Gly-36 to Gln-42, Gly-70 to Leu-77.	HCROB90R	
ID NO. 6556 as residues: Ser-15 to Ile-24, Asn-56 to Lys-67, Ser-80 to Lys-95. HCRPJ34R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6557 as residues: Val-5 to Gln-11. HCQBL95R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6558 as residues: Ser-4 to Pro-10, Glu-18 to Cys-23. HWLOR95 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6559 as residues: Ser-23 to Ala-28, Pro-64 to Glu-74, Ala-100 to Lys-106. HKCSI32R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6560 as residues: Ala-4 to Gln-14, Gly-36 to Gln-42, Gly-70 to Leu-77.		ID NO. 6555 as residues: Arg-63 to Gly-69.
HCRPJ34R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6557 as residues: Val-5 to Gln-11. HCQBL95R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6558 as residues: Ser-4 to Pro-10, Glu-18 to Cys-23. HWLOR95 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6559 as residues: Ser-23 to Ala-28, Pro-64 to Glu-74, Ala-100 to Lys-106. HKCSI32R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6560 as residues: Ala-4 to Gln-14, Gly-36 to Gln-42, Gly-70 to Leu-77.	HCRNI50R	
HCRPJ34R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6557 as residues: Val-5 to Gln-11. HCQBL95R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6558 as residues: Ser-4 to Pro-10, Glu-18 to Cys-23. HWLOR95 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6559 as residues: Ser-23 to Ala-28, Pro-64 to Glu-74, Ala-100 to Lys-106. HKCSI32R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6560 as residues: Ala-4 to Gln-14, Gly-36 to Gln-42, Gly-70 to Leu-77.		1 · · · · · · · · · · · · · · · · · · ·
ID NO. 6557 as residues: Val-5 to Gln-11. HCQBL95R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6558 as residues: Ser-4 to Pro-10, Glu-18 to Cys-23. HWLOR95 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6559 as residues: Ser-23 to Ala-28, Pro-64 to Glu-74, Ala-100 to Lys-106. HKCSI32R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6560 as residues: Ala-4 to Gln-14, Gly-36 to Gln-42, Gly-70 to Leu-77.		<u> </u>
HCQBL95R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6558 as residues: Ser-4 to Pro-10, Glu-18 to Cys-23. HWLOR95 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6559 as residues: Ser-23 to Ala-28, Pro-64 to Glu-74, Ala-100 to Lys-106. HKCSI32R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6560 as residues: Ala-4 to Gln-14, Gly-36 to Gln-42, Gly-70 to Leu-77.	HCRPJ34R	
ID NO. 6558 as residues: Ser-4 to Pro-10, Glu-18 to Cys-23. HWLOR95 R ID NO. 6559 as residues: Ser-23 to Ala-28, Pro-64 to Glu-74, Ala-100 to Lys-106. HKCSI32R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6560 as residues: Ala-4 to Gln-14, Gly-36 to Gln-42, Gly-70 to Leu-77.	HCORI 95R	
HWLOR95 R ID NO. 6559 as residues: Ser-23 to Ala-28, Pro-64 to Glu-74, Ala-100 to Lys-106. HKCSI32R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6560 as residues: Ala-4 to Gln-14, Gly-36 to Gln-42, Gly-70 to Leu-77.	1.0025556	
R ID NO. 6559 as residues: Ser-23 to Ala-28, Pro-64 to Glu-74, Ala-100 to Lys-106. HKCSI32R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6560 as residues: Ala-4 to Gln-14, Gly-36 to Gln-42, Gly-70 to Leu-77.	HWLOR95	
to Lys-106. HKCSI32R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6560 as residues: Ala-4 to Gln-14, Gly-36 to Gln-42, Gly-70 to Leu-77.		,
HKCS132R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6560 as residues: Ala-4 to Gln-14, Gly-36 to Gln-42, Gly-70 to Leu-77.		
ID NO. 6560 as residues: Ala-4 to Gln-14, Gly-36 to Gln-42, Gly-70 to Leu-77.	HKCSI32R	
Leu-77.		
	HBCJN86R	Preferred epitopes include those comprising a sequence shown in SEQ

	ID NO. 6565 as residues: Pro-6 to Tyr-17, Val-39 to Gln-45.
HWLMZ47	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6567 as residues: Ile-45 to Gly-50.
HCRPD88R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6569 as residues: Asn-15 to Phe-27, His-39 to Ser-44, Glu-49 to
	Ala-55.
HCQDC47R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6571 as residues: Asp-1 to Asn-7, Pro-22 to Ser-28, Leu-54 to
	Asn-59, Gly-95 to Arg-101.
H2CBR33R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6573 as residues: Ile-2 to Leu-8.
HWLXV36	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6574 as residues: Lys-14 to Gln-24, Pro-32 to Ile-40.
HWLRE24	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6575 as residues: Lys-20 to Gly-38, Val-42 to Thr-53, Ala-88 to
	Ala-99.
HWMBA27	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6576 as residues: Gly-35 to Glu-62.
HWMBK08	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6577 as residues: Asp-2 to Cys-8.
HCQCT96R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6582 as residues: Pro-1 to Glu-14.
HWLXR95	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6584 as residues: Lys-22 to Ser-33, Ala-39 to Glu-48, Lys-70 to
	Lys-75.
HEPAD45R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6586 as residues: Met-42 to Arg-53.
HCRNP41R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6587 as residues: Arg-25 to Asn-34, Lys-54 to Glu-60.
HCYBK83	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6588 as residues: Pro-1 to Ser-6.
HCRND59R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6589 as residues: Phe-88 to Pro-93, Thr-102 to Pro-113.
HCRMA15	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6592 as residues: Gly-4 to Lys-10, Gln-36 to Glu-41.
HCRMJ42R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6593 as residues: Gly-4 to Lys-10, Gln-36 to Glu-41.
HCRMO88	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6595 as residues: Gly-4 to Lys-10, Gln-36 to Glu-41, Phe-57 to
	Asn-62.
HCRNB87R	Preferred epitopes include those comprising a sequence shown in SEQ
1100	ID NO. 6596 as residues: Arg-17 to His-22.
HCRNL44R	Preferred epitopes include those comprising a sequence shown in SEQ
VVOD PVV 4 5	ID NO. 6598 as residues: Ser-2 to Ala-7.
HCRPK46R	Preferred epitopes include those comprising a sequence shown in SEQ
WODEN 105	ID NO. 6603 as residues: Tyr-3 to Gly-10, Ala-17 to Tyr-24.
HCRPK48R	Preferred epitopes include those comprising a sequence shown in SEQ
110500000	ID NO. 6604 as residues: Asn-1 to Arg-9.
HCRQG02R	Preferred epitopes include those comprising a sequence shown in SEQ

	ID NO. 6606 as residues: Tyr-1 to Gly-14.
HCRQM26	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6607 as residues: Tyr-1 to Gly-16.
HHMMA34	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6609 as residues: Gly-4 to Leu-11.
HHMMA44	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6610 as residues: Gly-4 to Lys-10, Gln-36 to Glu-41.
HHMMC42	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6611 as residues: Gly-4 to Lys-10.
ННММС86	Preferred epitopes include those comprising a sequence shown in SEQ
R .	ID NO. 6612 as residues: Gly-4 to Lys-10, Gln-36 to Pro-43.
ннммез8	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6614 as residues: Gly-4 to Lys-10, Gln-36 to Lys-43.
ННММЕ80	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6618 as residues: Gly-4 to Lys-10, Gln-36 to Lys-43.
HHMMF79	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6620 as residues: Val-2 to Gly-9.
HOCTA39R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6621 as residues: Lys-7 to Lys-19.
HULCG37R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6623 as residues: Ile-2 to Ser-15, Gln-30 to Asp-38.
HWLMQ27	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6629 as residues: Pro-16 to Tyr-23.
HWLMQ65	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6632 as residues: Gln-37 to Arg-42.
HWLNZ20	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6634 as residues: Pro-12 to Glu-21.
HWLNZ35	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6635 as residues: Pro-16 to Gly-35.
HWLNZ44	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6636 as residues: Pro-13 to Glu-22.
HWLOW58	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6639 as residues: Gly-4 to Lys-10.
HWMBS18	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6653 as residues: Pro-10 to Trp-21.
HCRPY45R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6662 as residues: Lys-7 to Lys-20, Gln-46 to Glu-51.
HHMMF44	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6667 as residues: Gly-8 to Leu-15, Gln-40 to Lys-48.
HTWEL13	Preferred epitopes include those comprising a sequence shown in SEQ
RA	ID NO. 6668 as residues: Cys-6 to Ser-12.
HCRMH46	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6674 as residues: Gln-19 to Glu-24.
HWLND45	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6676 as residues: Gly-4 to Lys-11.
HWLWG95	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6677 as residues: Arg-21 to Arg-36.
HCRQO33R	Preferred epitopes include those comprising a sequence shown in SEQ
L	ID NO. 6679 as residues: Pro-6 to Asp-21.

HCRMJ70R	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6682 as residues: Leu-18 to Asp-41.
HWLMM72	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6683 as residues: Asp-42 to Asn-47.
HCRMD32	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6684 as residues: Ala-17 to Lys-28, Glu-51 to Gln-56, Ser-64 to
	Lys-72.
HKAHM80	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6688 as residues: Lys-10 to Ala-17, Glu-27 to Leu-37, Met-74 to Lys-80, Pro-94 to Gln-108.
H2CBM60R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6689 as residues: His-23 to Arg-30, Asp-61 to Asn-73, Phe-89 to Gln-97.
HWLXR73	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6692 as residues: Arg-1 to Pro-11, Gly-16 to Gly-21, Gly-28 to Gly-43.
HWLOI59R	Preferred epitopes include those comprising a sequence shown in SEQ
II W E O IS S K	ID NO. 6693 as residues: Val-9 to Leu-20, Lys-44 to Pro-51.
HWLUX53	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6694 as residues: Gly-1 to Glu-10.
HARMO20	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6696 as residues: Arg-2 to Val-18.
HCQDM81 R	Preferred epitopes include those comprising a sequence shown in SEQ
HFIJB15R	ID NO. 6699 as residues: Arg-2 to Val-18. Preferred epitopes include those comprising a sequence shown in SEQ
III IJB13K	ID NO. 6703 as residues: Pro-26 to Gln-32.
HACCH14R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6704 as residues: Thr-1 to Tyr-7.
HCRPV08R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6708 as residues: Val-26 to Val-33, Phe-41 to Ser-55, Val-62 to Gly-72.
HWMBB77	Preferred epitopes include those comprising a sequence shown in SEQ
R ·	ID NO. 6709 as residues: Gln-7 to Leu-17, Lys-110 to Cys-116, Asn-133 to Asn-138.
HHEPL48R	Preferred epitopes include those comprising a sequence shown in SEQ
IIIIEI E46K	ID NO. 6710 as residues: Thr-2 to Met-11, Cys-15 to Pro-20, Asp-28 to
]	Ser-33, Lys-40 to Gly-45.
HCRPT53R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6711 as residues: Tyr-9 to Phe-14, Glu-30 to Lys-39.
HTXJU67R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6714 as residues: Ile-10 to Gln-15, Pro-22 to Asn-28.
HWMCL33	Preferred epitopes include those comprising a sequence shown in SEQ
R HCQCO67R	ID NO. 6715 as residues: Ala-83 to Ala-88.
HCQCOO/R	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6717 as residues: Cys-15 to Ser-30, Ser-39 to Met-45.
HWLV133R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6719 as residues: Arg-32 to Gly-38.
HWMBA55	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6720 as residues: Thr-1 to Gln-7, Thr-26 to Leu-36, Ala-86 to

	Asp-104, Ser-114 to Val-121.
HCRON89R	Preferred epitopes include those comprising a sequence shown in SEQ
HCKONOSK	ID NO. 6725 as residues: Ala-15 to Gly-22.
HLDDP53R	Preferred epitopes include those comprising a sequence shown in SEQ
HLUDFSSK	ID NO. 6727 as residues: Ala-25 to Asp-32.
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HWLME23	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6728 as residues: Ala-9 to Arg-15.
HWLVP88	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6729 as residues: Arg-21 to Ser-28, Gly-115 to Gln-142.
HWLMG29	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6739 as residues: Ser-16 to Lys-21, Pro-34 to Lys-41.
HCQCF55R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6740 as residues: Arg-1 to Arg-26, Ser-42 to Tyr-50, Glu-60 to
	Cys-69.
HWLWB88	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6742 as residues: Pro-6 to Glu-13.
HWLXR58	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6747 as residues: Glu-4 to Asp-12, Glu-19 to Lys-29, Ser-32 to
	Glu-40, Glu-51 to Thr-56, Ile-58 to Ser-79, Ser-86 to Glu-95.
HCYBO60R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6750 as residues: His-8 to Gly-18, Gly-26 to Pro-35, Pro-58 to
	Asp-64.
HE2BG62R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6751 as residues: Phe-10 to Tyr-15.
HCRMW12	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6752 as residues: Gly-21 to Asn-31, Cys-62 to Lys-68, Pro-76
	to Thr-81, Cys-105 to Arg-124, Lys-139 to Gln-145, Gly-151 to Gly-
	158.
HWLVF61	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6755 as residues: Tyr-12 to Ile-17, Pro-28 to Asn-33, Arg-45 to
	Asp-53.
HWMBP47	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6757 as residues: Val-1 to Val-10.
HWLQF89	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6760 as residues: Pro-8 to Pro-25, Asp-72 to Thr-78, Glu-81 to
}	Ser-87.
HWMCC54	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6763 as residues: Gln-66 to Ser-71, Ser-80 to Gly-92.
HCQAS76R	Preferred epitopes include those comprising a sequence shown in SEQ
l	ID NO. 6767 as residues: Thr-34 to Ser-40.
HKLRA71R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6768 as residues: Ile-1 to Ser-9.
HWMCJ58	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6769 as residues: Pro-10 to Arg-18.
HWLMJ20	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6770 as residues: Pro-56 to Trp-61.
HWLMU79	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6773 as residues: Trp-4 to Lys-11.
HWLNN06	Preferred epitopes include those comprising a sequence shown in SEQ
I LI AA I HAHATIA	

R	ID NO. 6775 as residues: Gln-27 to Ser-32, Trp-57 to Ser-65, Glu-72 to Ser-85, Lys-103 to Ser-117.
HWLMM42	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6776 as residues: Asn-43 to His-64.
HWMBC38	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6777 as residues: His-61 to Gly-68.
HWLVU11	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6778 as residues: Val-65 to Thr-74, Ser-84 to Asn-101.
HCQDW90	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6785 as residues: Arg-18 to Ser-24.
НСҮВМ34	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6786 as residues: Arg-22 to Ser-28.
HCYBM57	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6787 as residues: Arg-31 to Thr-38.
HCQCK49	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6789 as residues: Phe-14 to Ser-22.
HWLRQ41	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6792 as residues: Lys-13 to Asp-24.
HWLOC77	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6793 as residues: Phe-47 to Ser-52.
HDDNQ21	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6794 as residues: Tyr-6 to Gly-13, Asn-35 to Thr-42, Pro-47 to
	Glu-56.
HCQDA89	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6796 as residues: Leu-7 to Arg-13.
HCQCO43R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6797 as residues: Asp-18 to Arg-29.
HCQCG73R	Preferred epitopes include those comprising a sequence shown in SEQ
no Quo, sic	ID NO. 6798 as residues: Lys-12 to Asn-18, Glu-24 to Glu-31, Ile-40 to
	Ala-53, Pro-65 to Asp-75.
HWLQA92	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6799 as residues: Arg-10 to Ser-18, Pro-27 to Lys-36.
HCROM41	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6802 as residues: Ser-1 to Arg-9, Thr-40 to Trp-47, Ser-84 to
I K	Asp-95, Leu-113 to Asn-127, Pro-140 to Arg-151.
H2LAA02R	
112LAAU2K	Preferred epitopes include those comprising a sequence shown in SEQ
HCODI 130	ID NO. 6803 as residues: Ala-11 to Pro-20, Asn-39 to Val-46.
HCQDU29	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6805 as residues: Val-1 to Met-8.
HWMBJ73	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6811 as residues: Arg-41 to Glu-46.
HCRNO44R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6814 as residues: Lys-1 to Thr-6.
HSAMD89	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6818 as residues: Leu-32 to Glu-59, Lys-67 to Lys-89.
HCROE42R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6820 as residues: Phe-48 to Gly-56, Ile-60 to Glu-65, Pro-73 to
	Trp-80, Ser-100 to Lys-117, Lys-126 to Ser-138.
HCROE77R	Preferred epitopes include those comprising a sequence shown in SEQ

	ID NO. 6824 as residues: Asp-46 to Lys-51.
HOCTA19R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6827 as residues: Ser-3 to Ala-12, Gly-71 to Val-84.
HWLOM88	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6829 as residues: Glu-11 to Cys-17, Ala-26 to Trp-31, Ser-43 to
	Glu-55, Gly-127 to Ala-132.
H2CBI14R	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6830 as residues: Lys-21 to Lys-29.
HCRNI08R	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6831 as residues: Glu-10 to Val-16, Thr-59 to Ser-66, Asp-112
	to Ala-121, Pro-147 to Ala-157.
HFPBS29R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6832 as residues: Pro-22 to His-30.
HCQCB43R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6834 as residues: Asn-4 to Tyr-9.
HCQDB27R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6836 as residues: Asn-4 to Tyr-9.
HCQCR82R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6838 as residues: Glu-9 to Gly-17.
HWLWH33	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6842 as residues: Arg-10 to Arg-15, Val-25 to Gly-33, Pro-45
	to Asp-51.
HCYBJ83R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6843 as residues: Arg-1 to Gly-6, Arg-60 to Gly-65.
HWLRE17	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6844 as residues: Gln-34 to Gly-46, Gly-54 to Arg-61, Pro-67
	to Gly-82, Glu-91 to Asn-114.
HWLOM10	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6846 as residues: Glu-1 to Arg-11, Thr-18 to Ser-39, Ala-51 to
	Leu-56, Pro-69 to Gly-78, Glu-88 to Ala-93, Pro-114 to Lys-126, Leu-133 to Thr-141.
H2LBA48R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6848 as residues: Thr-13 to Thr-23.
HCRPZ16R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6849 as residues: Ala-14 to Cys-32, Lys-34 to Arg-40, Ser-46 to
	Trp-52, Arg-59 to Gly-64.
HKCSA80R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6850 as residues: Asn-39 to Gln-44.
HCRPH64R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6852 as residues: Arg-38 to Ser-46.
HDTBZ03R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6853 as residues: Lys-1 to Gly-28, Thr-50 to Leu-57, Glu-70 to
	Trp-90, Pro-93 to Asp-100.
HLYED39R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6854 as residues: Arg-2 to Thr-9.
HCQCB85R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6855 as residues: Gly-9 to Ser-14, Gln-26 to Gly-37.
HCRNF48R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6858 as residues: Glu-29 to Leu-34, Thr-40 to Pro-45, Ser-68 to

HWLQA11 Referred epitopes include those comprising a sequence shown in SEQ ID NO. 6860 as residues: His-60 to Cys-69. HWLXJ34R DFeferred epitopes include those comprising a sequence shown in SEQ ID NO. 6863 as residues: Arg-13 to Leu-22, Ser-25 to Glu-30, Leu-32 to Ala-43, Thr-49 to Pro-55, Ala-69 to Tyr-76, Pro-83 to Ser-91, Glu-104 to Ser-115. HCRQN67R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6865 as residues: Lys-1 to Ser-12, Arg-20 to Gln-25, Pro-80 to Arg-86. HCYBH30R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6866 as residues: Thr-19 to Lys-27. HCROE26R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6869 as residues: Arg-25 to Val-33, Ser-43 to Gly-48, Ala-54 to Gly-59. HOHBE57R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6870 as residues: Asp-1 to Gln-14, Thr-34 to Pro-40, Asn-42 to Asp-57, Ala-112 to Gly-117. HWMBB94 R DNO. 6873 as residues: Ser-53 to Val-62. HUVHA17 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6873 as residues: Glu-34 to Thr-41. HLTIJ91R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6874 as residues: Glu-34 to Thr-41. HCRMC40 R DNO. 6873 as residues: Glu-31 to Leu-21, Glu-42 to Gln-50. HCRMC40 R DNO. 6873 as residues: Glu-31 to Leu-21, Glu-42 to Gln-50. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6874 as residues: Ala-37 to Lys-42, Pro-55 to Asp-62. HWLQD31 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6883 as residues: Ala-37 to Lys-42, Pro-55 to Asp-62. HCSBE19R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6883 as residues: Ala-37 to Lys-42, Pro-55 to Asp-62. HYLQG37 R DNO. 6883 as residues: Ala-10 to Lys-16, Lys-19 to Val-27. HSAMB82 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6883 as residues: Ala-10 to Lys-16, Lys-19 to Val-27. Preferred epitopes include those c		Met-73.
HWLXJ34R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6863 as residues: Arg-13 to Leu-22, Ser-25 to Glu-30, Leu-32 to Ala-43, Thr-49 to Pro-55, Ala-69 to Tyr-76, Pro-83 to Ser-91, Glu-104 to Ser-115. HCRQN67R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6865 as residues: Lys-1 to Ser-12, Arg-20 to Gln-25, Pro-80 to Arg-86. HCYBH30R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6866 as residues: Thr-19 to Lys-27. HCROE26R HCROE26R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6869 as residues: Arg-25 to Val-33, Ser-43 to Gly-48, Ala-54 to Gly-59. HOHBE57R HOHBE57R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6870 as residues: Asp-1 to Gln-14, Thr-34 to Pro-40, Asn-42 to Asp-57, Ala-112 to Gly-117. HWMBB94 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6873 as residues: Ser-53 to Val-62. HUVHA17 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6873 as residues: Glu-34 to Thr-41. HLTIJ91R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6873 as residues: Glu-11 to Leu-21, Glu-42 to Gln-50. HCRMC40 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6875 as residues: Arg-37 to Val-48. HWLQD31 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6883 as residues: Arg-37 to Val-48. HWLQG37 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6883 as residues: Ala-37 to Lys-42, Pro-55 to Asp-62. HOSBE19R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6883 as residues: Ala-10 to Lys-16, Lys-19 to Val-27. HSAMB82 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6883 as residues: Ala-10 to Lys-16, Lys-19 to Val-27. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6893 as residues: Thr-5 to Thr-14. HFVLM205 Preferred epitopes include thos	HWLQA11	Preferred epitopes include those comprising a sequence shown in SEQ
D NO. 6863 as residues: Arg-13 to Leu-22, Ser-25 to Glu-30, Leu-32 to Ala-43, Thr-49 to Pro-55, Ala-69 to Tyr-76, Pro-83 to Ser-91, Glu-104 to Ser-115. HCRQN67R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6865 as residues: Lys-1 to Ser-12, Arg-20 to Gln-25, Pro-80 to Arg-86. HCYBH30R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6866 as residues: Thr-19 to Lys-27. HCROE26R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6869 as residues: Arg-25 to Val-33, Ser-43 to Gly-48, Ala-54 to Gly-59. HOHBE57R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6870 as residues: Asp-1 to Gln-14, Thr-34 to Pro-40, Asn-42 to Asp-57, Ala-112 to Gly-117. HWMBB94 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6872 as residues: Ser-53 to Val-62. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6873 as residues: Glu-34 to Thr-41. HLTIJ91R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6873 as residues: Glu-34 to Thr-41. HCRMC40 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6873 as residues: Glu-31 to Leu-21, Glu-42 to Gln-50. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6878 as residues: Ala-37 to Lys-42, Pro-55 to Asp-62. HOSBEJ9R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6888 as residues: Ala-37 to Lys-42, Pro-55 to Asp-62. HWLQG37 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6888 as residues: Ala-10 to Lys-16, Lys-19 to Val-27. HSAMB82 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6884 as residues: Ala-10 to Lys-16, Lys-19 to Val-27. HYLWE05 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6893 as residues: Gln-1 to Lys-21, Lys-19 to Val-27. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6894 as residues: Thr-5 to Thr-	R	ID NO. 6860 as residues: His-60 to Cys-69.
to Ala-43, Thr-49 to Pro-55, Āla-69 to Tyr-76, Pro-83 to Ser-91, Glu- 104 to Ser-115. HCRQN67R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6865 as residues: Lys-1 to Ser-12, Arg-20 to Gln-25, Pro-80 to Arg-86. HCYBH30R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6866 as residues: Thr-19 to Lys-27. HCROE26R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6869 as residues: Arg-25 to Val-33, Ser-43 to Gly-48, Ala-54 to Gly-59. HOHBE57R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6870 as residues: Asp-1 to Gln-14, Thr-34 to Pro-40, Asn-42 to Asp-57, Ala-112 to Gly-117. HWMBB94 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6872 as residues: Ser-53 to Val-62. HUVHA17 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6873 as residues: Glu-34 to Thr-41. HLTU91R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6874 as residues: Glu-11 to Leu-21, Glu-42 to Gln-50. HCRMC40 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6878 as residues: Arg-37 to Val-48. HWLQD31 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6878 as residues: Ala-37 to Lys-42, Pro-55 to Asp-62. HOSBE19R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6880 as residues: Ala-37 to Lys-42, Pro-55 to Nay-62. HSAMB82 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6883 as residues: Ala-10 to Lys-16, Lys-19 to Val-27. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6883 as residues: Thr-5 to Thr-14. HFVKA92R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6893 as residues: Thr-5 to Thr-14. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6893 as residues: Thr-5 to Thr-14. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6894 as resid	HWLXJ34R	Preferred epitopes include those comprising a sequence shown in SEQ
HCRQN67R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6865 as residues: Lys-1 to Ser-12, Arg-20 to Gln-25, Pro-80 to Arg-86. HCYBH30R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6866 as residues: Thr-19 to Lys-27. HCROE26R HCROE26R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6869 as residues: Arg-25 to Val-33, Ser-43 to Gly-48, Ala-54 to Gly-59. HOHBE57R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6870 as residues: Asp-1 to Gln-14, Thr-34 to Pro-40, Asn-42 to Asp-57, Ala-112 to Gly-117. HWMBB94 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6872 as residues: Ser-53 to Val-62. HUVHA17 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6873 as residues: Glu-34 to Thr-41. HLTIJ91R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6874 as residues: Glu-34 to Thr-41. HCROE40 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6874 as residues: Ala-37 to Val-42 to Gln-50. HCRMC40 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6875 as residues: Arg-37 to Val-48. HWLQD31 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6878 as residues: Arg-37 to Val-48. HWLQG37 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6880 as residues: Ala-37 to Lys-42, Pro-55 to Asp-62. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6882 as residues: Ala-10 to Lys-16, Lys-19 to Val-27. HSAMB82 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6884 as residues: Thr-5 to Thr-14. HFVKA92R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6893 as residues: Thr-5 to Thr-14. HFVKA92R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6894 as residues: Pho-1 to Glu-12, Gln-21 to Asp-28, Asp-30 to Pro-35. HWLNK27 Preferred epitopes incl		ID NO. 6863 as residues: Arg-13 to Leu-22, Ser-25 to Glu-30, Leu-32
HCRQN67R DNO. 6865 as residues: Lys-1 to Ser-12, Arg-20 to Gln-25, Pro-80 to Arg-86. HCYBH30R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6866 as residues: Thr-19 to Lys-27. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6869 as residues: Arg-25 to Val-33, Ser-43 to Gly-48, Ala-54 to Gly-59. HCROE26R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6870 as residues: Asp-1 to Gln-14, Thr-34 to Pro-40, Asn-42 to Asp-57, Ala-112 to Gly-117. HWMBB94 R DNO. 6870 as residues: Ser-53 to Val-62. HUVHA17 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6873 as residues: Glu-34 to Thr-41. HLTIJ91R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6873 as residues: Glu-34 to Thr-41. HCRMC40 R DNO. 6874 as residues: Glu-11 to Leu-21, Glu-42 to Gln-50. HCRMC40 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6875 as residues: Arg-37 to Val-48. HWLQD31 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6878 as residues: Ala-37 to Lys-42, Pro-55 to Asp-62. HOSBE19R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6880 as residues: Ala-37 to Lys-42, Pro-55 to Asp-62. HSAMB82 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6880 as residues: Ala-10 to Lys-16, Lys-19 to Val-27. HSAMB82 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6883 as residues: Ala-10 to Lys-16, Lys-19 to Val-27. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6883 as residues: Ala-10 to Lys-16, Lys-19 to Val-27. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6894 as residues: Asp-28 to Arg-34. HKLSA82R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6894 as residues: Phe-1 to Glu-12, Gln-21 to Asp-28, Asp-30 to Pro-35. HWLNK27 Preferred epitopes include those comprising a sequence shown in SEQ ID		to Ala-43, Thr-49 to Pro-55, Ala-69 to Tyr-76, Pro-83 to Ser-91, Glu-
DNO. 6865 as residues: Lys-1 to Ser-12, Arg-20 to Gln-25, Pro-80 to Arg-86. HCYBH30R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6866 as residues: Thr-19 to Lys-27. HCROE26R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6869 as residues: Arg-25 to Val-33, Ser-43 to Gly-48, Ala-54 to Gly-59. HOHBE57R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6870 as residues: Asp-1 to Gln-14, Thr-34 to Pro-40, Asn-42 to Asp-57, Ala-112 to Gly-117. HWMBB94 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6872 as residues: Ser-53 to Val-62. HUVHA17 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6873 as residues: Glu-34 to Thr-41. HLTIJ91R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6874 as residues: Glu-11 to Leu-21, Glu-42 to Gln-50. HCRMC40 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6878 as residues: Arg-37 to Val-48. HWLQD31 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6878 as residues: Arg-37 to Val-48. HWLQD31 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6878 as residues: Ala-37 to Lys-42, Pro-55 to Asp-62. HOSBE19R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6880 as residues: Ala-37 to Lys-42, Pro-55 to Asp-62. HXLQQ37 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6882 as residues: Ala-10 to Lys-16, Lys-19 to Val-27. HXLWB05 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6884 as residues: Ala-10 to Lys-16, Lys-19 to Val-27. HYVKA92R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6893 as residues: Asp-28 to Arg-34. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6894 as residues: Ala-10 to Lys-16, Lys-19 to Asp-28, Asp-30 to Pro-35. HYVKA92R Preferred epitopes include those comprising a sequence shown		104 to Ser-115.
Arg-86. HCYBH30R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6866 as residues: Thr-19 to Lys-27. HCROE26R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6869 as residues: Arg-25 to Val-33, Ser-43 to Gly-48, Ala-54 to Gly-59. HOHBE57R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6870 as residues: Asp-1 to Gln-14, Thr-34 to Pro-40, Asn-42 to Asp-57, Ala-112 to Gly-117. HWMBB94 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6872 as residues: Ser-53 to Val-62. HUVHA17 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6873 as residues: Glu-34 to Thr-41. HLTIJ91R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6874 as residues: Glu-11 to Leu-21, Glu-42 to Gln-50. HCRMC40 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6875 as residues: Arg-37 to Val-48. HWLQD31 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6878 as residues: Ala-37 to Lys-42, Pro-55 to Asp-62. HOSBE19R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6888 as residues: Ala-37 to Lys-42, Pro-55 to Asp-62. HWLQG37 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6882 as residues: Ala-10 to Lys-16, Lys-19 to Val-27. HSAMB82 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6883 as residues: Gln-1 to Arg-11. HFVKA92R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6894 as residues: Asp-25 to Arg-34. HKLSA82R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6894 as residues: Asp-28 to Arg-34. HKLSA82R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6894 as residues: Office of the pro-35. HWLNK27 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6894 as residues: Office of the pro-35. HCRN724R Preferred epitopes include those comprisin	HCRQN67R	Preferred epitopes include those comprising a sequence shown in SEQ
HCYBH30R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6866 as residues: Thr-19 to Lys-27. HCROE26R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6869 as residues: Arg-25 to Val-33, Ser-43 to Gly-48, Ala-54 to Gly-59. HOHBE57R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6870 as residues: Asp-1 to Gln-14, Thr-34 to Pro-40, Asn-42 to Asp-57, Ala-112 to Gly-117. HWMBB94 R ID NO. 6872 as residues: Ser-53 to Val-62. HUVHA17 R ID NO. 6873 as residues: Ser-53 to Val-62. HUTIJ91R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6873 as residues: Glu-34 to Thr-41. HLTIJ91R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6874 as residues: Glu-11 to Leu-21, Glu-42 to Gln-50. HCRMC40 R ID NO. 6875 as residues: Arg-37 to Val-48. HWLQD31 R ID NO. 6878 as residues: Arg-37 to Val-48. HWLQD31 R ID NO. 6878 as residues: Arg-37 to Lys-42, Pro-55 to Asp-62. HOSBE19R ID NO. 6880 as residues: Asp-25 to Ile-31. HWLQG37 R ID NO. 6880 as residues: Asp-25 to Ile-31. HWLQG37 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6888 as residues: Asp-25 to IR-31. HWLWG03 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6888 as residues: Asp-25 to IR-31. HWLWG03 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6888 as residues: Thr-5 to Thr-14. HFVKA92R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6893 as residues: Asp-28 to Arg-34. HKLSA82R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6896 as residues: Asp-28 to Arg-34. HKLSA82R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6896 as residues: Phe-1 to Glu-12, Gln-21 to Asp-28, Asp-30 to Pro-35. HWLNX27 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6896 as residues: Gln-2 to Trp-8. HCRNT24R Preferred epitopes include those comprising a sequence		ID NO. 6865 as residues: Lys-1 to Ser-12, Arg-20 to Gln-25, Pro-80 to
HCROE26R HCROE26R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6869 as residues: Arg-25 to Val-33, Ser-43 to Gly-48, Ala-54 to Gly-59. HOHBE57R ID NO. 6870 as residues: Asp-1 to Gln-14, Thr-34 to Pro-40, Asn-42 to Asp-57, Ala-112 to Gly-117. HWMBB94 R ID NO. 6872 as residues: Ser-53 to Val-62. HUVHA17 R ID NO. 6873 as residues: Ser-53 to Val-62. HUVHA17 R ID NO. 6873 as residues: Glu-34 to Thr-41. HLTIJ91R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6873 as residues: Glu-31 to Tu-41. HCRMC40 R ID NO. 6874 as residues: Glu-11 to Leu-21, Glu-42 to Gln-50. HCRMC40 R ID NO. 6875 as residues: Arg-37 to Val-48. HWLQD31 R ID NO. 6876 as residues: Ala-37 to Val-48. HWLQD31 R ID NO. 6880 as residues: Ala-37 to Lys-42, Pro-55 to Asp-62. HOSBE19R ID NO. 6880 as residues: Ala-37 to Lys-42, Pro-55 to Asp-62. HOSBE19R ID NO. 6880 as residues: Ala-10 to Lys-16, Lys-19 to Val-27. HSAMB82 R ID NO. 6883 as residues: Gln-1 to Arg-11. HWLWE05 R ID NO. 6884 as residues: Thr-5 to Thr-14. HFVKA92R ID NO. 6883 as residues: Asp-28 to Arg-34. HKLSA82R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6893 as residues: Asp-28 to Arg-34. HKLSA82R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6894 as residues: Gln-1 to Arg-11. HWLNK27 R ID NO. 6895 as residues: Asp-28 to Arg-34. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6896 as residues: Gln-2 to Trp-8. HCRNT24R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6896 as residues: Gln-2 to Trp-8. HCRNT24R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6897 as residues: Gln-2 to Trp-8. HCRNT24R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6898 as residues: Gln-2 to Trp-8. HCRNT24R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6898 as residues: Thr-2 to Lys-7, Lys-12 to Pro-21.		
HCROE26R ID NO. 6869 as residues: Arg-25 to Val-33, Ser-43 to Gly-48, Ala-54 to Gly-59. HOHBE57R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6870 as residues: Asp-1 to Gln-14, Thr-34 to Pro-40, Asn-42 to Asp-57, Ala-112 to Gly-117. HWMBB94 R ID NO. 6872 as residues: Ser-53 to Val-62. HUVHA17 R ID NO. 6873 as residues: Ser-53 to Val-62. HUVHA17 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6873 as residues: Glu-34 to Thr-41. HLTIJ91R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6874 as residues: Glu-34 to Thr-41. HCRMC40 R ID NO. 6875 as residues: Glu-11 to Leu-21, Glu-42 to Gln-50. HCRMC40 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6875 as residues: Arg-37 to Val-48. HWLQD31 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6878 as residues: Ala-37 to Lys-42, Pro-55 to Asp-62. HOSBE19R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6888 as residues: Ala-37 to Lys-42, Pro-55 to Asp-62. HWLQG37 R ID NO. 6882 as residues: Ala-10 to Lys-16, Lys-19 to Val-27. HSAMB82 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6882 as residues: Gln-1 to Arg-11. HWLWE05 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6884 as residues: Thr-5 to Thr-14. HFVKA92R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6894 as residues: Phe-1 to Glu-12, Gln-21 to Asp-28, Asp-30 to Pro-35. HWLNX27 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6894 as residues: Phe-1 to Glu-12, Gln-21 to Asp-28, Asp-30 to Pro-35. HWLNX27 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6896 as residues: Phe-1 to Glu-12, Gln-21 to Asp-28, Asp-30 to Pro-35. HWLNX27 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6896 as residues: Gln-2 to Trp-8. HCRNT24R Preferred epitopes include those comprising a sequence shown	HCYBH30R	Preferred epitopes include those comprising a sequence shown in SEQ
ID NO. 6869 as residues: Arg-25 to Val-33, Ser-43 to Gly-48, Ala-54 to Gly-59. HOHBE57R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6870 as residues: Asp-1 to Gln-14, Thr-34 to Pro-40, Asn-42 to Asp-57, Ala-112 to Gly-117. HWMBB94 R ID NO. 6872 as residues: Ser-53 to Val-62. HUVHA17 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6873 as residues: Glu-34 to Thr-41. HLTIJ91R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6874 as residues: Glu-11 to Leu-21, Glu-42 to Gln-50. HCRMC40 R ID NO. 6874 as residues: Arg-37 to Val-48. HWLQD31 R ID NO. 6875 as residues: Ala-37 to Lys-42, Pro-55 to Asp-62. HOSBE19R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6888 as residues: Ala-37 to Lys-42, Pro-55 to Asp-62. HWLQG37 R ID NO. 6880 as residues: Ala-10 to Lys-16, Lys-19 to Val-27. HSAMB82 R ID NO. 6883 as residues: Gln-1 to Arg-11. HWLWE05 R ID NO. 6884 as residues: Gln-1 to Arg-11. HWLWE05 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6884 as residues: Ala-10 to Lys-16, Lys-19 to Val-27. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6884 as residues: Ala-10 to Arg-11. HYKA92R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6894 as residues: Asp-28 to Arg-34. HKLSA82R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6894 as residues: Asp-28 to Arg-34. HKLSA82R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6896 as residues: Asp-28 to Arg-34. HKLSA82R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6896 as residues: Asp-21 to Trp-8. HCRNT24R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6896 as residues: Gln-2 to Trp-8. HCRNT24R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6898 as residues: Arg-13 to Thr-21, Ser-43 to Ala-49. Preferred epitopes include those comprising		ID NO. 6866 as residues: Thr-19 to Lys-27.
Gly-59. HOHBE57R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6870 as residues: Asp-1 to Gln-14, Thr-34 to Pro-40, Asn-42 to Asp-57, Ala-112 to Gly-117. HWMBB94 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6872 as residues: Ser-53 to Val-62. HUVHA17 R ID NO. 6873 as residues: Glu-34 to Thr-41. HLTIJ91R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6874 as residues: Glu-34 to Thr-41. HCRMC40 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6875 as residues: Ala-37 to Val-48. HWLQD31 R ID NO. 6875 as residues: Ala-37 to Val-48. HWLQD31 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6888 as residues: Ala-37 to Lys-42, Pro-55 to Asp-62. HOSBE19R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6880 as residues: Asp-25 to Ile-31. HWLQG37 R ID NO. 6883 as residues: Ala-10 to Lys-16, Lys-19 to Val-27. HSAMB82 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6883 as residues: Gln-1 to Arg-11. HWLWE05 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6884 as residues: Thr-5 to Thr-14. HFVKA92R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6893 as residues: Asp-28 to Arg-34. HKLSA82R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6894 as residues: Phe-1 to Glu-12, Gln-21 to Asp-28, Asp-30 to Pro-35. HWLNK27 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6896 as residues: Gln-2 to Trp-8. HCRNT24R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6896 as residues: Gln-2 to Trp-8. HCRNT24R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6898 as residues: Gln-2 to Trp-8. HCRNT24R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6898 as residues: Thr-2 to Lys-7, Lys-12 to Pro-21. HFCES53R Preferred epitopes include those comprisin	HCROE26R	
Gly-59. HOHBE57R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6870 as residues: Asp-1 to Gln-14, Thr-34 to Pro-40, Asn-42 to Asp-57, Ala-112 to Gly-117. HWMBB94 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6872 as residues: Ser-53 to Val-62. HUVHA17 R ID NO. 6873 as residues: Glu-34 to Thr-41. HLTIJ91R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6874 as residues: Glu-34 to Thr-41. HCRMC40 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6875 as residues: Ala-37 to Val-48. HWLQD31 R ID NO. 6875 as residues: Ala-37 to Val-48. HWLQD31 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6888 as residues: Ala-37 to Lys-42, Pro-55 to Asp-62. HOSBE19R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6880 as residues: Asp-25 to Ile-31. HWLQG37 R ID NO. 6883 as residues: Ala-10 to Lys-16, Lys-19 to Val-27. HSAMB82 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6883 as residues: Gln-1 to Arg-11. HWLWE05 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6884 as residues: Thr-5 to Thr-14. HFVKA92R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6893 as residues: Asp-28 to Arg-34. HKLSA82R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6894 as residues: Phe-1 to Glu-12, Gln-21 to Asp-28, Asp-30 to Pro-35. HWLNK27 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6896 as residues: Gln-2 to Trp-8. HCRNT24R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6896 as residues: Gln-2 to Trp-8. HCRNT24R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6898 as residues: Gln-2 to Trp-8. HCRNT24R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6898 as residues: Thr-2 to Lys-7, Lys-12 to Pro-21. HFCES53R Preferred epitopes include those comprisin		ID NO. 6869 as residues: Arg-25 to Val-33, Ser-43 to Gly-48, Ala-54 to
ID NO. 6870 as residues: Asp-1 to Gln-14, Thr-34 to Pro-40, Asn-42 to Asp-57, Ala-112 to Gly-117. HWMBB94 R DNO. 6872 as residues: Ser-53 to Val-62. HUVHA17 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6873 as residues: Glu-34 to Thr-41. HLTIJ91R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6874 as residues: Glu-11 to Leu-21, Glu-42 to Gln-50. HCRMC40 R DNO. 6874 as residues: Glu-11 to Leu-21, Glu-42 to Gln-50. HCRMC40 R DNO. 6875 as residues: Arg-37 to Val-48. HWLQD31 R DNO. 6878 as residues: Ala-37 to Lys-42, Pro-55 to Asp-62. HOSBE19R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6880 as residues: Asp-25 to Ile-31. HWLQG37 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6882 as residues: Ala-10 to Lys-16, Lys-19 to Val-27. HSAMB82 R DNO. 6883 as residues: Gln-1 to Arg-11. HWLWE05 R DNO. 6884 as residues: Thr-5 to Thr-14. HFVKA92R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6893 as residues: Thr-5 to Thr-14. HFVKA92R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6894 as residues: Thr-5 to Thr-14. HKLSA82R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6894 as residues: Gln-1 to Glu-12, Gln-21 to Asp-28, Asp-30 to Pro-35. HWLNK27 R DNO. 6896 as residues: Gln-2 to Trp-8. HCRNT24R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6896 as residues: Gln-2 to Trp-8. HCRNT24R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6897 as residues: Arg-13 to Thr-21, Ser-43 to Ala-49. HCQAW95 R DNO. 6898 as residues: Thr-2 to Lys-7, Lys-12 to Pro-21. HFCES53R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6898 as residues: Thr-2 to Lys-7, Lys-12 to Pro-21.		
Asp-57, Ala-112 to Gly-117. HWMBB94 R Dreferred epitopes include those comprising a sequence shown in SEQ ID NO. 6872 as residues: Ser-53 to Val-62. HUVHA17 R DNO. 6873 as residues: Glu-34 to Thr-41. HLTIJ91R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6874 as residues: Glu-11 to Leu-21, Glu-42 to Gln-50. HCRMC40 R DNO. 6875 as residues: Ala-37 to Val-48. HWLQD31 R DNO. 6875 as residues: Ala-37 to Lys-42, Pro-55 to Asp-62. HOSBE19R DNO. 6876 as residues: Ala-37 to Lys-42, Pro-55 to Asp-62. HOSBE19R DNO. 6880 as residues: Ala-37 to Lys-42, Pro-55 to Asp-62. HOSBE19R DNO. 6882 as residues: Ala-37 to Lys-16, Lys-19 to Val-27. HSAMB82 R DNO. 6882 as residues: Ala-10 to Lys-16, Lys-19 to Val-27. HSAMB82 R DNO. 6883 as residues: Gln-1 to Arg-11. HWLWE05 R DNO. 6884 as residues: Gln-1 to Arg-11. HWLWE05 R DNO. 6883 as residues: Thr-5 to Thr-14. HFVKA92R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6893 as residues: Asp-28 to Arg-34. HKLSA82R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6893 as residues: Asp-28 to Arg-34. HKLSA82R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6893 as residues: Asp-28 to Arg-34. HKLSA82R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6894 as residues: Phe-1 to Glu-12, Gln-21 to Asp-28, Asp-30 to Pro-35. HWLNK27 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6896 as residues: Gln-2 to Trp-8. HCRNT24R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6896 as residues: Arg-13 to Thr-21, Ser-43 to Ala-49. HCQAW95 R DNO. 6893 as residues: Thr-2 to Lys-7, Lys-12 to Pro-21. HFCES53R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6893 as residues: Thr-2 to Lys-7, Lys-12 to Pro-21.	HOHBE57R	Preferred epitopes include those comprising a sequence shown in SEQ
HWMBB94 R DNO. 6872 as residues: Ser-53 to Val-62. HUVHA17 R DNO. 6873 as residues: Ser-53 to Val-62. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6873 as residues: Glu-34 to Thr-41. HLTIJ91R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6874 as residues: Glu-11 to Leu-21, Glu-42 to Gln-50. HCRMC40 R DNO. 6875 as residues: Arg-37 to Val-48. HWLQD31 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6875 as residues: Ala-37 to Lys-42, Pro-55 to Asp-62. HOSBE19R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6880 as residues: Asp-25 to Ile-31. HWLQG37 R DNO. 6882 as residues: Ala-10 to Lys-16, Lys-19 to Val-27. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6883 as residues: Gln-1 to Arg-11. HWLWE05 R DNO. 6883 as residues: Thr-5 to Thr-14. HFVKA92R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6893 as residues: Asp-28 to Arg-34. HKLSA82R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6894 as residues: Phe-1 to Glu-12, Gln-21 to Asp-28, Asp-30 to Pro-35. HWLNK27 R DNO. 6896 as residues: Gln-2 to Trp-8. HCRNT24R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6896 as residues: Gln-2 to Trp-8. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6896 as residues: Gln-2 to Trp-8. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6897 as residues: Asp-13 to Thr-21, Ser-43 to Ala-49. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6898 as residues: Arg-13 to Thr-21, Ser-43 to Ala-49. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6898 as residues: Thr-2 to Lys-7, Lys-12 to Pro-21.		ID NO. 6870 as residues: Asp-1 to Gln-14, Thr-34 to Pro-40, Asn-42 to
R ID NO. 6872 as residues: Ser-53 to Val-62. HUVHA17 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6873 as residues: Glu-34 to Thr-41. HLTIJ91R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6874 as residues: Glu-11 to Leu-21, Glu-42 to Gln-50. HCRMC40 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6875 as residues: Arg-37 to Val-48. HWLQD31 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6878 as residues: Ala-37 to Lys-42, Pro-55 to Asp-62. HOSBE19R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6880 as residues: Asp-25 to Ile-31. HWLQG37 R ID NO. 6880 as residues: Ala-10 to Lys-16, Lys-19 to Val-27. HSAMB82 R ID NO. 6882 as residues: Gln-1 to Arg-11. HWLWE05 R ID NO. 6884 as residues: Gln-1 to Arg-11. HYLWE05 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6884 as residues: Asp-28 to Arg-34. HFVKA92R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6893 as residues: Asp-28 to Arg-34. HKLSA82R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6894 as residues: Asp-28 to Arg-34. PREFERRED PREFERRE		
HUVHA17 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6873 as residues: Glu-34 to Thr-41. HLTIJ91R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6874 as residues: Glu-11 to Leu-21, Glu-42 to Gln-50. HCRMC40 R ID NO. 6875 as residues: Arg-37 to Val-48. HWLQD31 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6878 as residues: Ala-37 to Lys-42, Pro-55 to Asp-62. HOSBE19R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6880 as residues: Asp-25 to Ile-31. HWLQG37 R ID NO. 6882 as residues: Ala-10 to Lys-16, Lys-19 to Val-27. HSAMB82 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6883 as residues: Gln-1 to Arg-11. HWLWE05 R ID NO. 6884 as residues: Thr-5 to Thr-14. HFVKA92R HKLSA82R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6893 as residues: Asp-28 to Arg-34. HKLSA82R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6894 as residues: Phe-1 to Glu-12, Gln-21 to Asp-28, Asp-30 to Pro-35. HWLNK27 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6894 as residues: Phe-1 to Glu-12, Gln-21 to Asp-28, Asp-30 to Pro-35. HWLNK27 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6896 as residues: Gln-2 to Trp-8. HCRNT24R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6897 as residues: Arg-13 to Thr-21, Ser-43 to Ala-49. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6898 as residues: Arg-13 to Thr-21, Ser-43 to Ala-49. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6898 as residues: Thr-2 to Lys-7, Lys-12 to Pro-21.	HWMBB94	Preferred epitopes include those comprising a sequence shown in SEQ
HLTIJ91R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6874 as residues: Glu-11 to Leu-21, Glu-42 to Gln-50. HCRMC40 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6875 as residues: Arg-37 to Val-48. HWLQD31 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6878 as residues: Ala-37 to Lys-42, Pro-55 to Asp-62. HOSBE19R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6880 as residues: Asp-25 to Ile-31. HWLQG37 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6882 as residues: Ala-10 to Lys-16, Lys-19 to Val-27. HSAMB82 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6883 as residues: Gln-1 to Arg-11. HWLWE05 R ID NO. 6884 as residues: Thr-5 to Thr-14. HFVKA92R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6893 as residues: Asp-28 to Arg-34. HKLSA82R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6894 as residues: Phe-1 to Glu-12, Gln-21 to Asp-28, Asp-30 to Pro-35. HWLNK27 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6894 as residues: Gln-2 to Trp-8. HCRNT24R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6896 as residues: Gln-2 to Trp-8. HCRNT24R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6897 as residues: Arg-13 to Thr-21, Ser-43 to Ala-49. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6898 as residues: Thr-2 to Lys-7, Lys-12 to Pro-21.		
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HCRMC40 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6875 as residues: Arg-37 to Val-48. HWLQD31 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6878 as residues: Ala-37 to Lys-42, Pro-55 to Asp-62. HOSBE19R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6880 as residues: Asp-25 to Ile-31. HWLQG37 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6882 as residues: Ala-10 to Lys-16, Lys-19 to Val-27. HSAMB82 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6883 as residues: Gln-1 to Arg-11. HWLWE05 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6884 as residues: Thr-5 to Thr-14. HFVKA92R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6893 as residues: Asp-28 to Arg-34. HKLSA82R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6894 as residues: Phe-1 to Glu-12, Gln-21 to Asp-28, Asp-30 to Pro-35. HWLNK27 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6896 as residues: Gln-2 to Trp-8. HCRNT24R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6897 as residues: Arg-13 to Thr-21, Ser-43 to Ala-49. HCQAW95 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6898 as residues: Thr-2 to Lys-7, Lys-12 to Pro-21. HFCES53R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6898 as residues: Thr-2 to Lys-7, Lys-12 to Pro-21.	R	
HCRMC40 R DNO. 6875 as residues: Arg-37 to Val-48. HWLQD31 R DNO. 6878 as residues: Arg-37 to Val-48. HWLQD31 R DNO. 6878 as residues: Ala-37 to Lys-42, Pro-55 to Asp-62. HOSBE19R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6880 as residues: Asp-25 to Ile-31. HWLQG37 R DNO. 6880 as residues: Asp-25 to Ile-31. HWLQG37 R DNO. 6882 as residues: Ala-10 to Lys-16, Lys-19 to Val-27. HSAMB82 R DNO. 6883 as residues: Gln-1 to Arg-11. HWLWE05 R DNO. 6884 as residues: Thr-5 to Thr-14. HFVKA92R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6893 as residues: Asp-28 to Arg-34. HKLSA82R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6894 as residues: Phe-1 to Glu-12, Gln-21 to Asp-28, Asp-30 to Pro-35. HWLNK27 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6896 as residues: Gln-2 to Trp-8. HCRNT24R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6897 as residues: Gln-2 to Trp-8. HCRNT24R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6897 as residues: Gln-2 to Trp-8. HCRNT24R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6897 as residues: Arg-13 to Thr-21, Ser-43 to Ala-49. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6898 as residues: Thr-2 to Lys-7, Lys-12 to Pro-21. HFCES53R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6898 as residues: Thr-2 to Lys-7, Lys-12 to Pro-21.	HLTIJ91R	Preferred epitopes include those comprising a sequence shown in SEQ
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		ID NO. 6898 as residues: Thr-2 to Lys-7, Lys-12 to Pro-21.
ID NO. 6902 as residues: Thr-12 to Leu-18.	HFCES53R	
	L	ID NO. 6902 as residues: Thr-12 to Leu-18.

HCQCQ84R	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6903 as residues: Gly-1 to Ala-10.
HWMBC92	Preferred epitopes include those comprising a sequence shown in SEQ
R R	ID NO. 6905 as residues: Leu-49 to Asn-62, Pro-65 to Leu-84.
HWLQQ35	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6908 as residues: Arg-6 to Ala-19, Asn-26 to Thr-50, Phe-57 to
	Ser-62, Asp-68 to Glu-96, Ser-102 to Gly-137.
HCRNZ02R	Preferred epitopes include those comprising a sequence shown in SEQ
11014.00211	ID NO. 6911 as residues: Asn-1 to Lys-9, Cys-51 to Ala-65, Thr-74 to
	Arg-86.
HCQDW65	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6916 as residues: Lys-19 to Ser-27.
HCQDN27	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6918 as residues: Glu-6 to Gln-21.
HCQCI92R	Preferred epitopes include those comprising a sequence shown in SEQ
11000	ID NO. 6919 as residues: Pro-19 to Lys-40.
HCROT79R	Preferred epitopes include those comprising a sequence shown in SEQ
nck01/9k	ID NO. 6922 as residues: Gly-12 to Glu-18.
11001 1070	
H2CAA07R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6923 as residues: Glu-8 to Ala-16, Tyr-25 to Trp-32.
H2LAD20R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6924 as residues: Ser-1 to Leu-6, Ser-22 to Leu-31.
HWLQZ32	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6925 as residues: Pro-1 to Leu-7, Gly-49 to Gly-69, Glu-100 to
	Ala-106.
HCRQK79	Preferred epitopes include those comprising a sequence shown in SEQ
Ř	ID NO. 6929 as residues: Lys-7 to Gly-14, Ala-31 to Gly-37.
HCQAD53	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6930 as residues: Thr-1 to Thr-13.
HKCUD58	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6931 as residues: Ser-21 to Cys-28.
HCRNR93R	Preferred epitopes include those comprising a sequence shown in SEQ
IICKINK95K	ID NO. 6932 as residues: Lys-54 to Leu-64.
HWLQH13	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6933 as residues: Asp-12 to Ser-19, Leu-52 to Gln-57, Leu-79
1	to Glu-86, Asn-97 to Phe-109, Gln-134 to Asn-142, Arg-151 to Gly-156.
HICHOSOR	Preferred epitopes include those comprising a sequence shown in SEQ
H2CBQ60R	ID NO. 6934 as residues: Ala-23 to Asp-32, Thr-42 to Gly-47, Pro-59
1101 111110	to Glu-67, Phe-77 to Ser-84.
H2LAW43	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6935 as residues: Thr-3 to Ser-12.
HWLVJ22R	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 6936 as residues: Gln-7 to Ser-23, Pro-63 to Lys-86.
H2CAA28R	Preferred epitopes include those comprising a sequence shown in SEQ
112CAA26R	ID NO. 6938 as residues: Glu-17 to Cys-22.
H2CAA36R	Preferred epitopes include those comprising a sequence shown in SEQ
III CANSON	ID NO. 6939 as residues: Asp-1 to Arg-9.
H2CBG84R	Preferred epitopes include those comprising a sequence shown in SEQ
HZCDU04K	ID NO. 6941 as residues: Gly-13 to Leu-20.
ī	ID NO. 0941 as residues: Gly-13 to Leu-20.

H2CBJ35R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6942 as residues: Val-3 to Ala-11, Ala-38 to Leu-51, Ser-53 to
via con via i b	Pro-70, Gln-88 to Gly-94, Ser-106 to Ser-113.
H2CBK71R	Preferred epitopes include those comprising a sequence shown in SEQ
3100001000	ID NO. 6944 as residues: Pro-18 to Pro-24, Arg-31 to Thr-41.
H2CBN87R	Preferred epitopes include those comprising a sequence shown in SEQ
H2CBP73R	ID NO. 6945 as residues: Asp-1 to Ser-6. Preferred epitopes include those comprising a sequence shown in SEQ
nzCBP/3R	ID NO. 6946 as residues: Ala-2 to Ser-9, Pro-40 to Gly-54.
H2CBS94R	Preferred epitopes include those comprising a sequence shown in SEQ
IIZOBOJ-IK	ID NO. 6947 as residues: Gly-39 to Gln-45.
H2CBV81R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6949 as residues: Arg-1 to Trp-8.
H2CBW73	Preferred epitopes include those comprising a sequence shown in SEQ
RB	ID NO. 6950 as residues: Trp-1 to Ser-8, Pro-17 to Glu-27, Gln-41 to
	Val-54, Asp-65 to Pro-76.
H2LAZ29R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6953 as residues: Asp-8 to Gly-18, Ala-21 to Arg-26, Glu-31 to
	Lys-36, Ser-61 to Gly-66.
H2LAZ92R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6954 as residues: His-10 to Phe-16, Thr-64 to Arg-79.
H2LBB20R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6956 as residues: Pro-17 to Arg-29, Gly-49 to Ala-62, Gly-70 to
HBAHC91R	Lys-81. Preferred epitopes include those comprising a sequence shown in SEQ
IDANCSIK	ID NO. 6960 as residues: Gln-21 to Ala-27.
HCEOM04	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6962 as residues: Thr-2 to Lys-11.
HCFOE14R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6963 as residues: Glu-20 to Tyr-25, Phe-43 to Glu-48.
HCHOX67	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6964 as residues: Ser-16 to His-21, Ala-29 to Thr-35.
HCQAB27R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6965 as residues: Lys-1 to Val-13.
HCQAB44R	Preferred epitopes include those comprising a sequence shown in SEQ
77.00 : 0.00	ID NO. 6968 as residues: Thr-19 to Thr-31.
HCQAB53R	Preferred epitopes include those comprising a sequence shown in SEQ
HCO A COST	ID NO. 6969 as residues: Ile-34 to His-39.
HCQAC03R	Preferred epitopes include those comprising a sequence shown in SEQ
HCOAD62	ID NO. 6970 as residues: Ser-51 to Gly-60. Preferred epitopes include those comprising a sequence shown in SEQ
HCQAD62 R	ID NO. 6976 as residues: Ala-1 to Val-8, Arg-24 to Gly-36.
HCQAE39R	Preferred epitopes include those comprising a sequence shown in SEQ
1.0Q. ILSSK	ID NO. 6981 as residues: Thr-3 to Arg-19.
HCQAG32	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6986 as residues: Arg-1 to Tyr-6.
HCQAI15R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 6994 as residues: Gly-1 to Ala-8.
HCQAK16	Preferred epitopes include those comprising a sequence shown in SEQ

R	ID NO. 6998 as residues: Gly-1 to Ser-9.
HCQAK17	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 6999 as residues: Ala-1 to Arg-7.
HCQAL71R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7001 as residues: Val-2 to His-12.
HCQAM57	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 7004 as residues: Arg-1 to Thr-8.
HCQAN95	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 7012 as residues: Phe-11 to Ser-17, Leu-42 to Gly-47.
HCQAR63R	Preferred epitopes include those comprising a sequence shown in SEQ
l	ID NO. 7016 as residues: Thr-5 to Arg-11.
HCQAS25R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7019 as residues: His-4 to His-10.
HCQAT12R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7025 as residues: Trp-2 to Gly-9.
HCQAV66	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 7032 as residues: Gly-1 to Ser-8.
HCQAW40	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 7036 as residues: His-1 to Ile-26, Leu-30 to Ser-37, Ala-59 to
	Leu-66.
HCQBA47R	Preferred epitopes include those comprising a sequence shown in SEQ
`	ID NO. 7038 as residues: Ser-8 to Arg-14.
HCQBE19R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7043 as residues: Glu-25 to Ser-30.
HCQBL61R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7049 as residues: Arg-38 to Asn-43.
HCQBM58	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 7052 as residues: Gln-7 to Glu-16.
HCQCC50R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7067 as residues: Arg-1 to Gly-8, Pro-11 to Asn-21, Gln-28 to
	Lys-36.
HCQCD10R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7072 as residues: Ser-33 to Tyr-42, Val-51 to Ser-56.
HCQCD46R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7073 as residues: Arg-14 to Thr-21.
HCQCE46R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7079 as residues: Ala-2 to Asp-10.
HCQCE83R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7085 as residues: Arg-14 to Thr-20.
HCQCF77R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7092 as residues: Lys-8 to Asn-19.
HCQCH16R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7104 as residues: Leu-31 to Thr-37, Gly-54 to Glu-61.
HCQCH47R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7108 as residues: Pro-13 to Glu-18.
HCQCJ42R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7126 as residues: Glu-1 to Gly-13.
HCQCJ51R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7129 as residues: Pro-8 to Asn-18, Gln-25 to Val-30.

HCQCJ77R	Preferred epitopes include those comprising a sequence shown in SEQ
11000000	ID NO. 7132 as residues: Asn-1 to Thr-6.
HCQCJ89R	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 7134 as residues: Phe-16 to Asn-27.
HCQCK81	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 7145 as residues: Glu-15 to Glu-20.
HCQCK90	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 7146 as residues: Pro-2 to Thr-10.
HCQCL01R	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 7147 as residues: Ser-10 to Gly-15.
HCQCL05R	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 7148 as residues: Thr-24 to Thr-33.
HCQCL14R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7151 as residues: Arg-3 to Gly-13.
HCQCL48R	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 7159 as residues: Ala-1 to Thr-13.
HCQCL51R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7160 as residues: Pro-9 to Asn-19.
HCQCL55R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7162 as residues: Pro-8 to Asn-18.
HCQCL65R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7165 as residues: Lys-1 to Gly-6, Glu-8 to Arg-13.
HCQCL78R	Preferred epitopes include those comprising a sequence shown in SEQ
110001 500	ID NO. 7169 as residues: Lys-15 to Asn-23.
HCQCL79R	Preferred epitopes include those comprising a sequence shown in SEQ
UCOCO20D	ID NO. 7170 as residues: Pro-1 to Pro-8, Pro-17 to Asp-44.
HCQCO30R	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 7174 as residues: Ala-17 to Asn-28.
HCQCO53R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7175 as residues: Asn-1 to Gly-11, Gly-16 to Arg-22.
HCQCO66R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7177 as residues: Phe-2 to Asn-11.
HCQCO79R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7178 as residues: Arg-1 to Arg-7.
HCQCP19R	Preferred epitopes include those comprising a sequence shown in SEQ
II.CO CD20D	ID NO. 7183 as residues: Arg-8 to Met-13, Leu-16 to Leu-24.
HCQCP30R	Preferred epitopes include those comprising a sequence shown in SEQ
HCQCP89R	ID NO. 7186 as residues: Lys-1 to His-7.
IICQCF69K	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 7193 as residues: Leu-42 to Ser-47.
HCQCR44R	Preferred epitopes include those comprising a sequence shown in SEQ
1100011111	ID NO. 7198 as residues: Lys-34 to Asn-40.
HCQCT38R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7200 as residues: Arg-18 to Arg-26.
HCQCU08R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7204 as residues: Lys-3 to Trp-8.
HCQCU57R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7208 as residues: Lys-1 to Lys-10.
HCQCU67R	Preferred epitopes include those comprising a sequence shown in SEQ

	ID NO. 7210 as residues: Phe-5 to Leu-13.
HCQCV50	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 7215 as residues: Thr-8 to Lys-14, Glu-38 to Thr-50, Arg-56 to
	Asp-62.
HCQCV91	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 7218 as residues: Lys-1 to Phe-11.
HCQCX90	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 7225 as residues: Leu-5 to Tyr-11.
HCQDA28	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 7228 as residues: Glu-48 to Lys-57.
HCQDA36	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 7229 as residues: Met-6 to Ser-14, Ser-24 to Lys-29.
HCQDA66	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 7232 as residues: Ala-10 to Thr-15, Arg-20 to Glu-34.
HCQDB17R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7234 as residues: Ala-2 to Gly-15, Cys-20 to Asn-29, Gln-35 to
	Lys-41, Phe-47 to Lys-59.
HCQDB41R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7237 as residues: Gly-1 to Ala-8.
HCQDB49R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7239 as residues: Phe-8 to Gly-13, Pro-16 to Asn-26, Gln-33 to
	Thr-38.
HCQDB52R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7240 as residues: Leu-13 to Ser-20.
HCQDB54R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7241 as residues: Pro-5 to Trp-17.
HCQDC12R	Preferred epitopes include those comprising a sequence shown in SEQ
HCOPPAG	ID NO. 7245 as residues: Glu-8 to Asn-13, Arg-16 to Ala-28.
HCQDD35	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 7255 as residues: Asn-26 to Tyr-32.
HCQDE68R	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 7269 as residues: Pro-8 to Asn-18, Leu-27 to Cys-33.
HCQDF44R	Preferred epitopes include those comprising a sequence shown in SEQ
ncQDF44K	ID NO. 7271 as residues: Ser-6 to Val-15.
HCQDF69R	Preferred epitopes include those comprising a sequence shown in SEQ
TICODIOSK	ID NO. 7274 as residues: Ser-19 to Arg-25.
HCQDG40	Preferred epitopes include those comprising a sequence shown in SEQ
R R	ID NO. 7278 as residues: Asn-2 to Val-8, Phe-25 to Leu-30.
HCQDG71	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 7280 as residues: Lys-8 to Phe-13.
HCQDG80	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 7281 as residues: Ser-4 to Tyr-10.
HCQDH18	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 7283 as residues: Asn-31 to Ser-37.
HCQDH60	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 7288 as residues: Pro-9 to Asn-19, Gln-26 to Ser-34.
HCQDJ22R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7306 as residues: Gly-9 to Asn-14.
HCQDK50	Preferred epitopes include those comprising a sequence shown in SEQ
	

R	ID NO. 7320 as residues: Lys-38 to Asp-43.
HCQDK58	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 7322 as residues: Lys-1 to Trp-6.
HCQDL36R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7327 as residues: Arg-12 to Ser-20.
HCQDL57R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7331 as residues: Ser-25 to Asp-32.
HCQDL96R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7333 as residues: Ser-8 to Ala-18.
HCQDM58	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 7338 as residues: Phe-5 to Ala-10.
HCQDN78	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 7342 as residues: Asn-1 to Gly-6, Pro-9 to Ser-14.
HCQDP14R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7351 as residues: Gly-1 to Tyr-13.
HCQDQ80	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 7359 as residues: Pro-34 to Ser-40.
HCQDS61R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7372 as residues: Ile-17 to Val-24.
HCQDU60	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 7387 as residues: Pro-9 to Asn-19.
HCQDU94	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 7390 as residues: Pro-7 to His-19.
HCQDV44	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 7393 as residues: Thr-19 to Thr-26, Ala-38 to Arg-43.
HCRMB19	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 7419 as residues: His-23 to Gln-29.
HCRMB44	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 7420 as residues: Ser-1 to Ser-8.
HCRMB82	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 7422 as residues: Pro-1 to Ser-9.
HCRMD33	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 7429 as residues: Pro-14 to Asn-21, Pro-23 to Asn-34.
HCRMD57	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 7430 as residues: Arg-14 to Ser-30.
HCRMD77	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 7431 as residues: Asn-4 to Asn-10.
HCRMF07	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 7436 as residues: Arg-1 to Gly-10, Glu-16 to Gln-21.
HCRMF33	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 7439 as residues: Pro-3 to Thr-8.
HCRMF93	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 7447 as residues: Leu-2 to Arg-9, Glu-23 to His-34.
HCRMG20	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 7449 as residues: Ser-15 to His-22.
HCRMI33R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7457 as residues: Phe-4 to Ala-10.
HCRMI60R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7460 as residues: Glu-21 to Gly-41, Ala-75 to Gly-80.

HCRMJ54R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7463 as residues: Pro-13 to Phe-23.
HCRMJ81R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7465 as residues: Phe-15 to Phe-24, Asn-63 to Ala-69, Leu-80
	to Pro-85.
HCRMP32	Preferred epitopes include those comprising a sequence shown in SEQ
RA	ID NO. 7472 as residues: Arg-5 to Glu-14, Arg-31 to Gly-36.
HCRMS48	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 7477 as residues: Arg-42 to Lys-50.
HCRMT03	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 7480 as residues: Phe-5 to Ser-13.
HCRMU21	Preferred epitopes include those comprising a sequence shown in SEQ
R HCRMW62	ID NO. 7483 as residues: Ser-20 to Glu-28.
	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 7497 as residues: Cys-53 to Ser-60.
R HCRMY29	Preferred epitopes include those comprising a sequence shown in SEQ
R R	ID NO. 7502 as residues: Arg-1 to Thr-6.
HCRMZ36	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 7505 as residues: Pro-7 to Ser-27.
HCRMZ71	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 7507 as residues: Gly-1 to Cys-7, Thr-33 to Lys-38.
HCRMZ92	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 7508 as residues: Gly-45 to Ile-56.
HCRNB85R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7519 as residues: His-1 to Arg-9.
HCRNC23R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7520 as residues: Lys-3 to Arg-11, Pro-19 to Gly-24, Ser-74 to
	Trp-79.
HCRNE15R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7527 as residues: Arg-7 to Ser-12.
HCRNE60R	Preferred epitopes include those comprising a sequence shown in SEQ
HODATEOLD	ID NO. 7532 as residues: Glu-1 to Ser-11.
HCRNF01R	Preferred epitopes include those comprising a sequence shown in SEQ
HCRNH02R	ID NO. 7533 as residues: Gly-46 to Thr-52. Preferred epitopes include those comprising a sequence shown in SEQ
11CKINIU2K	ID NO. 7538 as residues: Asn-46 to Gly-57.
HCRNI71R	Preferred epitopes include those comprising a sequence shown in SEQ
licidii/iik	ID NO. 7540 as residues: Lys-1 to Trp-10.
HCRNJ25R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7541 as residues: Asp-10 to His-16, Arg-24 to Trp-29, Lys-40
	to Phe-46, Leu-83 to Trp-90, Pro-92 to His-97.
HCRNK40	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 7543 as residues: 1le-49 to Asn-55, Ser-69 to His-79.
HCRNK94	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 7544 as residues: Met-34 to Pro-48.
HCRNL38R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7546 as residues: Ser-11 to Ser-16, Ala-52 to Glu-60.
HCRNL55R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7548 as residues: Thr-7 to Thr-15.

HCRNM50	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 7553 as residues: Ser-18 to Asn-26.
HCRNO49R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7559 as residues: Gly-24 to Arg-36, Pro-57 to Arg-65.
HCRNV70	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 7571 as residues: Asn-1 to Lys-6, Ser-14 to Gly-26.
HCRNW29	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 7573 as residues: Gly-23 to Ser-28.
HCRNX03	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 7577 as residues: Arg-1 to Glu-9.
HCRNZ22R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7581 as residues: Leu-24 to Asp-32.
HCROE81R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7592 as residues: Gly-1 to Thr-8.
HCROE89R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7593 as residues: Gly-13 to His-18.
HCROF67R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7595 as residues: Lys-1 to Asn-19, Thr-61 to Ala-68.
HCROG58R	Preferred epitopes include those comprising a sequence shown in SEQ
110000000	ID NO. 7599 as residues: Pro-44 to Gly-49.
HCROG62R	Preferred epitopes include those comprising a sequence shown in SEQ
VICE CYTOOD	ID NO. 7600 as residues: Ser-19 to Pro-26.
HCROH29R	Preferred epitopes include those comprising a sequence shown in SEQ
HCDOISSD	ID NO. 7602 as residues: Thr-34 to Ser-40, Arg-102 to Trp-109.
HCROJ88R	Preferred epitopes include those comprising a sequence shown in SEQ
HCROK42	ID NO. 7613 as residues: Arg-26 to Gly-33, Arg-39 to Arg-60.
R R	Preferred epitopes include those comprising a sequence shown in SEQ
HCROK47	ID NO. 7619 as residues: Arg-20 to Met-28. Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 7620 as residues: Arg-8 to Pro-13.
HCROM53	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 7631 as residues: Val-11 to Gln-17, Pro-41 to Thr-47, Arg-66 to
	Glu-75.
HCROM56	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 7632 as residues: Arg-12 to Asn-17, Cys-26 to Gln-36.
HCRON01R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7636 as residues: Asp-4 to Thr-10.
HCRON04R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7637 as residues: Thr-1 to Pro-9.
HCRON70R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7641 as residues: Gly-1 to Arg-12.
HCROO46R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7643 as residues: Gln-47 to Ser-58.
HCROQ92R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7653 as residues: Ser-16 to Ser-28.
HCROR76R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7656 as residues: Ser-6 to Gly-11.
HCROS08R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7658 as residues: Asn-23 to Asn-29.
HCROS08R	Preferred epitopes include those comprising a sequence shown in SEQ

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HCROT15R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7662 as residues: Pro-26 to Lys-39, Asn-42 to Asn-49.
HCROT84R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7666 as residues: Pro-22 to Gly-28, Gly-37 to Lys-44.
HCROW69	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 7674 as residues: Arg-1 to Gly-8, Leu-19 to Pro-25.
HCROX18	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 7676 as residues: Gly-1 to Arg-9.
HCROX38	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 7678 as residues: Gly-3 to Val-9.
HCROZ45R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7683 as residues: Thr-1 to Gln-9, Thr-19 to Ser-31, Pro-36 to
	Glu-42, Leu-53 to Ala-63, Asn-92 to Gly-98, Leu-124 to Leu-131.
HCRPA19R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7689 as residues: Phe-62 to His-68.
HCRPA91R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7692 as residues: Gln-15 to Asn-26.
HCRPC30R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7695 as residues: Val-1 to Gly-6.
HCRPC56R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7698 as residues: Arg-1 to Glu-11, Val-27 to Val-35.
HCRPC58R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7699 as residues: Ala-4 to Thr-9.
HCRPE32R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7705 as residues: Asp-1 to Asp-18, Ser-41 to Arg-52.
HCRPE74R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7706 as residues: Met-6 to Gln-17.
HCRPF62R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7708 as residues: Cys-16 to Lys-33.
HCRPG28R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7715 as residues: Pro-26 to Ser-32.
HCRPG37R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7716 as residues: Arg-3 to Arg-9.
HCRPH31R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7719 as residues: Pro-35 to Gly-40.
HCRPH50R	Preferred epitopes include those comprising a sequence shown in SEQ
A	ID NO. 7720 as residues: Pro-2 to His-8.
HCRPH58R	Preferred epitopes include those comprising a sequence shown in SEQ
A	ID NO. 7721 as residues: Arg-14 to Val-19.
HCRPJ68R	Preferred epitopes include those comprising a sequence shown in SEQ
A	ID NO. 7727 as residues: Trp-29 to Asn-42.
HCRPL63R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7733 as residues: Ser-10 to Leu-21, Phe-31 to Lys-36, Ala-54 to
	Leu-67.
HCRPL79R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7734 as residues: Arg-1 to Leu-6.
HCRPM51	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 7737 as residues: Gly-14 to Thr-19, Gly-42 to Trp-48, Asp-63
	to Ala-71.

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HCRPN29R	Preferred epitopes include those comprising a sequence shown in SEQ
neid Nesic	ID NO. 7740 as residues: Lys-7 to Cys-12.
HCRPN49R	Preferred epitopes include those comprising a sequence shown in SEQ
IICIG NASIC	ID NO. 7742 as residues: Ser-6 to Thr-11, Pro-14 to His-28, Pro-34 to
	Asp-42, Pro-51 to Thr-60.
HCRPN73R	Preferred epitopes include those comprising a sequence shown in SEQ
Held Will	ID NO. 7743 as residues: Asn-16 to Ala-21.
HCRPO31R	Preferred epitopes include those comprising a sequence shown in SEQ
meid offic	ID NO. 7746 as residues: Gly-25 to Arg-30.
HCRPQ72R	Preferred epitopes include those comprising a sequence shown in SEQ
11010 Q/210	ID NO. 7754 as residues: Pro-12 to Ser-17, Trp-30 to Ala-35, Gln-49 to
}	Gln-55.
HCRPR62R	Preferred epitopes include those comprising a sequence shown in SEQ
1	ID NO. 7757 as residues: Cys-14 to His-20.
HCRPR70R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7758 as residues: Arg-16 to His-24.
HCRPR91R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7759 as residues: Tyr-1 to Ile-6, Gln-16 to Asp-24.
HCRPT82R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7767 as residues: Lys-1 to Lys-7.
HCRPU09R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7769 as residues: Phe-20 to Thr-25.
HCRPV91R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7775 as residues: Glu-19 to Ala-31, Glu-52 to Thr-82, Leu-104
	to Gln-110, Arg-125 to Arg-130.
HCRPX71R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7779 as residues: Pro-5 to Ala-11.
HCRPY01R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7780 as residues: Glu-1 to Gly-10, Ala-23 to Phe-33, Gln-59 to
	Ser-64.
HCRPY91R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7782 as residues: His-9 to Thr-17, Thr-25 to His-31.
HCRQB75R	Preferred epitopes include those comprising a sequence shown in SEQ
HOD COACD	ID NO. 7785 as residues: Arg-11 to Gly-23.
HCRQC36R	Preferred epitopes include those comprising a sequence shown in SEQ
HODODOOD	ID NO. 7786 as residues: Arg-53 to Arg-60.
HCRQD29R	Preferred epitopes include those comprising a sequence shown in SEQ
HCD CD 47D	ID NO. 7788 as residues: Pro-7 to Ala-15, Ser-32 to Lys-40.
HCRQD47R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7790 as residues: Ser-57 to Arg-64, Glu-71 to Gly-84, Arg-95
HCRQJ26R	to Trp-100.
TCKQJ20K	Preferred epitopes include those comprising a sequence shown in SEQ
HCRQL13R	ID NO. 7804 as residues: Asn-1 to Gly-9.
11CKQL13K	Preferred epitopes include those comprising a sequence shown in SEQ
HCRQL65R	ID NO. 7808 as residues: Glu-22 to Gly-27.
nckylosk	Preferred epitopes include those comprising a sequence shown in SEQ
HCPOM27	ID NO. 7809 as residues: Arg-6 to Thr-11.
HCRQM37 R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7810 as residues: Ala-42 to Pro-47, Pro-59 to Ser-66, Leu-79 to

	Arg-84, Gly-114 to Thr-119, Pro-132 to Gly-139.
HCRQM58	Preferred epitopes include those comprising a sequence shown in SEQ
Ř	ID NO. 7812 as residues: Glu-1 to Thr-7, Leu-12 to Asn-18.
HCRQM59	Preferred epitopes include those comprising a sequence shown in SEQ
Ř	ID NO. 7813 as residues: Glu-6 to Gly-13, Pro-64 to Ala-70.
HCYBA36R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7818 as residues: Tyr-40 to Ser-48.
HCYBD19R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7820 as residues: Ala-18 to Glu-26, Lys-39 to Glu-44, Phe-50
	to Ser-55.
HCYBE34R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7822 as residues: Glu-27 to Pro-34, Ser-49 to Gln-54, Ser-56 to
	Thr-62, Asp-102 to Lys-107, Gly-113 to Glu-119.
HCYBH89R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7826 as residues: Pro-33 to Pro-47.
HCYBH93R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7827 as residues: Ser-11 to Thr-19, Arg-59 to Asp-65.
HDPPE11R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7830 as residues: Pro-1 to Ala-14, Pro-44 to Gly-51.
HDTDS96R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7835 as residues: Ser-17 to Pro-22.
HE8AE77R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7840 as residues: Ile-3 to Asn-9.
HEONL43R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7842 as residues: Arg-1 to Val-10.
HFKHA60R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7845 as residues: Pro-13 to Arg-18, Phe-27 to Glu-37, Ala-45 to
	Leu-53, Gln-61 to Glu-69, Ser-75 to Ser-82, Gln-84 to Gly-94, Ala-96 to
	Pro-112.
HFRBW76	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 7847 as residues: Thr-2 to Gly-13.
HGBBA17R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7849 as residues: Asp-16 to Asn-22.
HHEQA63R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7853 as residues: Thr-13 to Ser-19, Ile-52 to Thr-59.
HHEWA82	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 7854 as residues: Cys-10 to Glu-15.
HHMMA39	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 7856 as residues: Arg-15 to Pro-21.
HHMMB13	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 7861 as residues: Glu-35 to Val-42.
HHMME20	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 7870 as residues: Thr-11 to Ala-17.
HJMBH59R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 7881 as residues: Ser-8 to Phe-24.
HKCSB18R	Preferred epitopes include those comprising a sequence shown in SEQ
1	Treferred epitopes include those comprising a sequence shown in 520
	ID NO. 7885 as residues: Arg-12 to Lys-19.
HKCSF11R	· · ·

HKCSJ63R	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 7892 as residues: Pro-6 to Gly-12.
HKCTB80R	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 7898 as residues: Ser-7 to Val-13, Arg-54 to Pro-62.
HKCTD27R	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 7900 as residues: Thr-9 to Gly-16.
HKLRA55R	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 7904 as residues: Arg-41 to Arg-47.
HKLSB04R	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 7912 as residues: Ser-27 to Leu-36, Glu-45 to Gly-52.
HKLSB05R	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 7913 as residues: Asn-1 to Phe-7, Val-15 to Met-20.
HKLSB41R	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 7914 as residues: Phe-13 to Ala-27, Gly-70 to Glu-77.
HKLSB76R	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 7915 as residues: Phe-1 to Gln-10.
HKLSC29R	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 7917 as residues: Ala-4 to Ser-12.
HKLSD79R	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 7922 as residues: Ser-8 to Gly-15.
HKLSD93R	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 7923 as residues: Gly-11 to Gly-17.
HNBTH48R	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 7928 as residues: Thr-7 to Ser-13.
HNTCO26R	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 7931 as residues: Arg-1 to Lys-10, Asn-18 to Thr-28.
HOCTA23R	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 7933 as residues: Phe-17 to Gly-22, Thr-40 to Val-47, Pro-58 to Gly-72, Pro-92 to Trp-109.
HOCTB19R	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 7936 as residues: Gln-13 to Ser-34.
HOCTB32R	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 7937 as residues: Arg-1 to Lys-8, Phe-30 to Lys-35.
HOCTC38R	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 7941 as residues: Ser-6 to Ser-14, Val-16 to Gln-23, Gly-39 to Ser-45, Thr-52 to Ser-58.
HOCTD35R	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 7946 as residues: Cys-2 to Val-7.
HOCTE12R	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 7950 as residues: Asn-1 to Val-6, Pro-22 to Phe-29.
HOCTF43R	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 7953 as residues: Asp-22 to Gly-27, Arg-35 to Pro-43, Asp-63 to Ser-68.
HOHAS78R	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 7959 as residues: Ala-1 to Cys-20, Arg-29 to Ser-37, Leu-48 to Phe-54.
HOSNW54 R	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 7961 as residues: Pro-1 to Asp-8, Asn-28 to Ser-33.
HPCRD42R	Preferred epitopes include those comprising a sequence shown in SEQ

	ID NO. 7963 as residues: Arg-1 to Glu-6, Arg-52 to Arg-57.
HPFCN76R	Preferred epitopes include those comprising a sequence shown in SEQ
III I CIV/OIC	ID NO. 7965 as residues: Ser-1 to Cys-16, Pro-30 to Asp-40.
HPJBZ88R	Preferred epitopes include those comprising a sequence shown in SEQ
III JDZ66K	ID NO. 7966 as residues: Pro-17 to Gly-27, Gly-30 to His-36, Phe-44 to
	Gly-54, Pro-56 to Ala-61.
HSIFC66R	Preferred epitopes include those comprising a sequence shown in SEQ
HSH COOK	ID NO. 7968 as residues: Glu-8 to Asn-13, Arg-16 to Thr-29.
HSOBF88R	Preferred epitopes include those comprising a sequence shown in SEQ
113ODF 66K	ID NO. 7969 as residues: Asp-1 to Tyr-8.
HSODE15R	Preferred epitopes include those comprising a sequence shown in SEQ
HSODETSK	ID NO. 7970 as residues: Leu-8 to Ser-15, Gly-21 to Ser-27.
HTXRF56R	Preferred epitopes include those comprising a sequence shown in SEQ
III AIG SOR	ID NO. 7974 as residues: Glu-1 to Arg-6, Ala-14 to Gly-27, Arg-31 to
	His-37.
HTYND19	Preferred epitopes include those comprising a sequence shown in SEQ
RA	ID NO. 7975 as residues: Glu-1 to Thr-15, Val-21 to Leu-27, Ser-37 to
	Arg-58, Met-82 to Asn-91.
HWLMA60	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 7981 as residues: Leu-10 to Arg-16.
HWLMB42	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 7984 as residues: Arg-24 to Arg-41.
HWLMC65	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 7985 as residues: Phe-18 to Trp-23.
HWLMC79	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 7986 as residues: Thr-30 to Thr-39.
HWLME59	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 7989 as residues: Asp-26 to Cys-32.
HWLME69	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 7990 as residues: Arg-11 to Gly-17.
HWLME71	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 7991 as residues: Gln-1 to Gly-6.
HWLMG12	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 7994 as residues: Asn-1 to Gly-10.
HWLMG15	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 7995 as residues: Pro-10 to Thr-16, Arg-39 to Gly-44.
HWLMG57	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8000 as residues: Ser-7 to Gly-17, Asn-35 to His-46.
HWLMG84	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8002 as residues: Ser-3 to Ala-23, Pro-25 to Gly-31, Ala-59 to Gly-80, Pro-83 to His-91, Gly-99 to Gly-110, Pro-112 to Trp-123.
HWLMH50	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8006 as residues: Ile-2 to Gln-7, Glu-21 to Gly-27.
HWLMJ80	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8010 as residues: Leu-65 to Thr-80.
HWLMK20	Preferred epitopes include those comprising a sequence shown in SEQ
R R	ID NO. 8011 as residues: Ser-8 to Pro-19.
HWLMK25	Preferred epitopes include those comprising a sequence shown in SEQ
R R	ID NO. 8012 as residues: Lys-1 to Ser-6.
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HWLMK31	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8013 as residues: Arg-1 to Trp-10, Arg-15 to Gly-24.
HWLMK62	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8014 as residues: Gly-1 to Ala-10, Pro-42 to Pro-53.
HWLMM68	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8015 as residues: His-10 to Asn-16.
HWLMQ01	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8023 as residues: Asn-3 to Lys-12.
HWLMR23	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8025 as residues: Gly-9 to Lys-17.
HWLMR69	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8026 as residues: Asp-6 to Glu-13, Leu-63 to Gln-70.
HWLMS31	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8027 as residues: Pro-2 to Leu-7.
HWLMT64	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8030 as residues: Asp-1 to Gln-6.
HWLMU26	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8033 as residues: Pro-21 to Val-26, Val-28 to Val-37, Ser-44 to
	Tyr-49, Phe-53 to Leu-65.
HWLMV60	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8036 as residues: Ser-27 to Glu-39, Leu-43 to Gln-48.
HWLNH76	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8050 as residues: Cys-8 to His-24, Ser-36 to Arg-44.
HWLNL41	Preferred epitopes include those comprising a sequence shown in SEQ
r R	ID NO. 8055 as residues: Pro-1 to Glu-22, Ala-31 to Asp-39, Glu-65 to
HWLNP65	Pro-72.
R	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8061 as residues: Val-12 to Trp-17, Ile-22 to Ser-28.
HWLNR26	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8063 as residues: Glu-10 to Gly-28.
HWLNY40	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8080 as residues: Pro-1 to Arg-18.
HWLOA09	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8083 as residues: Tyr-13 to Phe-18, Gln-22 to Tyr-27, Pro-74 to
	Met-81.
HWLOC65	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8087 as residues: Arg-41 to Asn-50.
HWLOF46	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8090 as residues: Arg-11 to Val-19, Thr-28 to Ala-39.
HWLOI17R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 8097 as residues: His-21 to Gly-29.
HWLOJ19R	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 8102 as residues: Ser-20 to Leu-37.
HWLOK12	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8105 as residues: Arg-24 to Asn-29.
HWLOK45	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8107 as residues: His-20 to Pro-26.
HWLON66	Description of anitary include these contributes are also in SEO
R	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8112 as residues: Phe-1 to Gln-11.

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HWLON71	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8113 as residues: Ala-1 to Tyr-8.
HWLOQ52	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8115 as residues: Cys-2 to Asn-8.
HWLOR15	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8117 as residues: Asp-1 to Gly-10, Thr-53 to Asp-59.
HWLOR65	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8118 as residues: Gly-16 to Gln-26, Gly-31 to Lys-37.
HWLOX29	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8132 as residues: Ser-16 to Ser-22.
HWLOY73	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8135 as residues: Pro-1 to Val-11, Pro-13 to Gln-20, Pro-39 to
	Pro-46, Gln-51 to Ala-73.
HWLOZ87	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8137 as residues: Gln-20 to Ser-27, Gln-42 to Ser-48.
HWLQA28	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8140 as residues: Lys-40 to Asn-55.
HWLQD30	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8147 as residues: Pro-6 to Pro-13, Gly-19 to Lys-39.
HWLQD40	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8148 as residues: Pro-14 to Asn-19, Glu-51 to Asn-57, Ser-67
WWW ODAG	to Pro-75.
HWLQD46	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8151 as residues: Gly-28 to Leu-33. Preferred epitopes include those comprising a sequence shown in SEQ
HWLQD89 R	ID NO. 8152 as residues: Lys-2 to Lys-7.
HWLQH32	Preferred epitopes include those comprising a sequence shown in SEQ
R R	ID NO. 8164 as residues: Asn-19 to Thr-27.
HWLQH58	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8165 as residues: Pro-45 to Asp-52.
HWLQM69	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8169 as residues: Glu-6 to Pro-12.
HWLQP18	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8175 as residues: Ser-2 to Ala-11.
HWLQQ83	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8177 as residues: Ser-26 to Gly-37, Pro-44 to Ser-50.
HWLQR90	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8178 as residues: Gln-1 to Trp-9, Val-17 to Glu-22.
HWLQT52	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8182 as residues: Gly-1 to Ser-10, Arg-16 to Met-22, Ser-24 to
	Trp-29, Gly-37 to Arg-44, Gly-52 to Ser-59, Arg-67 to Ser-85, Thr-107
	to Gly-114.
HWLQU50	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8185 as residues: Tyr-26 to Cys-34, Thr-45 to Asn-50.
HWLRB15	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8191 as residues: Leu-8 to His-14, Ser-17 to Trp-31, Thr-44 to
	Gln-50, Ala-53 to Ala-61, Thr-72 to Ala-90, Val-116 to Leu-123.
HWLRE01	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8194 as residues: Ser-9 to Asn-19, Asn-34 to Cys-41.

HWLRV63 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8218 as residues: Glu-15 to Cys-26, Arg-34 to Ille-58. HWLUG53 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8222 as residues: Asn-17 to Lys-27. HWLUH72 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8225 as residues: Asp-58 to Cys-72, Gln-81 to Glu-89. HWLUJ19R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8231 as residues: Ser-49 to Ser-55. HWLUL47 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8236 as residues: Lys-18 to Lys-24. HWLUUG5 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8237 as residues: Asp-1 to His-8. HWLUQ87 R ID NO. 8236 as residues: Cys-34 to Arg-41. HWLUUR41 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8256 as residues: Ser-24 to Trp-30. HWLUWB1 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8256 as residues: Ser-24 to Trp-30. HWLUV67 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8261 as residues: Pro-5 to Arg-13. HWLUZ07 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8269 as residues: Glu-1 to Gly-8. HWLVD26 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8259 as residues: Glu-1 to Gly-8. HWLVD74 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8278 as residues: Arg-1 to Asp-16. HWLVD74 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8278 as residues: Arg-54 to His-62. HWLV918 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8278 as residues: Arg-54 to His-62. HWLV918 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8290 as residues: Arg-54 to His-62. HWLVW60 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8290 as residues: Arg-54 to His-62. HWLVW10 Preferred epitopes include those	HWLRO35	Preferred epitopes include those comprising a sequence shown in SEQ
HWLUG53 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8225 as residues: Asp-58 to Cys-72, Gln-81 to Glu-89. HWLUJ19R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8225 as residues: Asp-58 to Cys-72, Gln-81 to Glu-89. HWLUJ19R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8231 as residues: Ser-49 to Ser-55. HWLUJL47 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8231 as residues: Lys-18 to Lys-24. HWLUJL47 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8236 as residues: Lys-18 to Lys-24. HWLUQ65 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8237 as residues: Cys-34 to Arg-41. HWLUQ87 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8254 as residues: Ser-24 to Trp-30. HWLUW88 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8256 as residues: Pro-9 to Gly-20. HWLUV67 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8261 as residues: Pro-9 to Gly-20. HWLUV07 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8264 as residues: Pro-5 to Arg-13. HWLVD06 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8269 as residues: Glu-1 to Gly-8. HWLVD16 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8279 as residues: Arg-11 to Asp-16. HWLVD27 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8279 as residues: Arg-11 to Asp-16. HWLVD18 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8279 as residues: Arg-14 to His-62. HWLV514 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8290 as residues: Arg-54 to His-62. HWLV734 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8290 as residues: Arg-54 to His-62. HWLV74 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8290 as	R	ID NO. 8206 as residues: Ile-20 to Thr-29, Lys-39 to Ala-46.
HWLUG53 R D NO. 8222 as residues: Asn-17 to Lys-27. R D NO. 8225 as residues: Asn-17 to Lys-27. R D NO. 8225 as residues: Asp-58 to Cys-72, Gln-81 to Glu-89. HWLUJ19R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8236 as residues: Sap-58 to Cys-72, Gln-81 to Glu-89. HWLUL47 R D NO. 8231 as residues: Lys-18 to Lys-24. HWLUL47 R D NO. 8236 as residues: Lys-18 to Lys-24. HWLUL65 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8237 as residues: Sap-1 to His-8. HWLUQ87 R HWLUQ87 R HWLUQ87 R R HWLUQ87 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8236 as residues: Cys-34 to Arg-41. HWLUU81 HWLUU81 R HWLUW81 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8256 as residues: Cys-34 to Arg-41. HWLUU88 R HWLUU089 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8256 as residues: Pro-9 to Gly-20. HWLUU070 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8264 as residues: Pro-9 to Gly-20. HWLUU707 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8269 as residues: Pro-5 to Arg-13. HWLVD207 R ID NO. 8269 as residues: Glu-1 to Gly-8. HWLVD40 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8275 as residues: Arg-11 to Asp-16. HWLVD74 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8278 as residues: Arg-11 to Asp-16. HWLVD74 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8278 as residues: Arg-11 to Thr-10. HWLVE21 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8282 as residues: Arg-14 to His-62. HWLVJ84 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8282 as residues: Arg-54 to His-62. HWLVJ84 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8290 as residues: Arg-54 to His-62. HWLVJ84 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8293 as residues: Arg-5	HWLRV63	Preferred epitopes include those comprising a sequence shown in SEQ
R ID NO. 8222 as residues: Asn-17 to Lys-27. HWLUH72 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8225 as residues: Asp-58 to Cys-72, Gln-81 to Glu-89. HWLUJ19R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8231 as residues: Ser-49 to Ser-55. HWLUL47 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8236 as residues: Lys-18 to Lys-24. HWLUJL65 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8236 as residues: Asp-1 to His-8. HWLUQ87 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8236 as residues: Cys-34 to Arg-41. HWLUW81 R HWLUW81 R HWLUW84 R ID NO. 8256 as residues: Cys-34 to Arg-41. HWLUU88 R ID NO. 8256 as residues: Ser-24 to Trp-30. HWLUU88 R ID NO. 8261 as residues: Pro-9 to Gly-20. R ID NO. 8264 as residues: Pro-9 to Gly-20. R ID NO. 8264 as residues: Pro-9 to Gly-20. R HWLUZ07 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8264 as residues: Pro-5 to Arg-13. HWLUZ07 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8269 as residues: Glu-1 to Gly-8. HWLVD64 R HWLVD74 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8275 as residues: Arg-11 to Asp-16. HWLVD74 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8278 as residues: His-1 to Thr-10. HWLVE21 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8278 as residues: His-1 to Thr-10. HWLVE14 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8279 as residues: His-1 to Thr-10. HWLVF34 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8289 as residues: His-1 to Thr-10. HWLV54 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8290 as residues: Arg-14 to His-62. HWLV744 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8290 as residues: Arg-14 to His-62. HWLVM62 Preferred epito	R	ID NO. 8218 as residues: Glu-15 to Cys-26, Arg-34 to Ile-58.
HWLUH72 R D NO. 8225 as residues: Asp-58 to Cys-72, Gln-81 to Glu-89. HWLUJ19R Preferred epitopes include those comprising a sequence shown in SEQ D NO. 8231 as residues: Ser-49 to Ser-55. HWLUL47 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8236 as residues: Lys-18 to Lys-24. HWLUL65 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8237 as residues: Lys-18 to Lys-24. HWLUQ87 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8236 as residues: Cys-34 to Arg-41. HWLUR41 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8254 as residues: Cys-34 to Arg-41. HWLUU88 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8256 as residues: Ser-24 to Trp-30. HWLUV67 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8261 as residues: Pro-9 to Gly-20. HWLUV67 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8264 as residues: Pro-9 to Arg-13. HWLUZ07 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8269 as residues: Pro-5 to Arg-13. HWLVD26 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8278 as residues: Arg-11 to Asp-16. HWLVD74 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8278 as residues: His-1 to Thr-10. HWLVE21 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8279 as residues: Arg-54 to His-62. HWLVF34 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8282 as residues: Arg-54 to His-62. HWLVJ15R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8289 as residues: Arg-54 to His-62. Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8290 as residues: Arg-10 Gly-33, Ser-35 to Phe-63. HWLV8A Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8293 as residues: Asr-2 to Gly-33, Ser-35 to Phe-63. HWLVM62 Preferred epitopes includ	HWLUG53	Preferred epitopes include those comprising a sequence shown in SEQ
HWLUJ19R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8231 as residues: Ser-49 to Ser-55. HWLUL47 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8236 as residues: Lys-18 to Lys-24. HWLUL65 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8237 as residues: Lys-18 to Lys-24. HWLUQ87 R ID NO. 8237 as residues: Asp-1 to His-8. HWLUQ87 R ID NO. 8254 as residues: Cys-34 to Arg-41. HWLUUR41 R IP referred epitopes include those comprising a sequence shown in SEQ ID NO. 8254 as residues: Cys-34 to Arg-41. HWLUUR41 R ID NO. 8256 as residues: Ser-24 to Trp-30. HWLUU88 R ID NO. 8261 as residues: Pro-9 to Gly-20. HWLUV67 R ID NO. 8264 as residues: Pro-9 to Gly-20. HWLUV707 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8264 as residues: Pro-5 to Arg-13. HWLVD20 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8269 as residues: Glu-1 to Gly-8. HWLVD26 R ID NO. 8275 as residues: Arg-11 to Asp-16. HWLVD74 R IP Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8275 as residues: Arg-11 to Thr-10. HWLVE21 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8278 as residues: His-1 to Thr-10. HWLVF34 R ID NO. 8278 as residues: His-1 to Thr-10. HWLVF34 R ID NO. 8279 as residues: Leu-33 to Glu-40, Lys-52 to Lys-62. HWLVJ34R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8289 as residues: Arg-54 to His-62. HWLVJ84R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8289 as residues: Arg-54 to His-62. HWLVB4 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8292 as residues: Arg-54 to His-62. HWLVM62 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8293 as residues: Arg-54 to His-62. HWLVM05 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8295 as residues: Arg-1 to Thr-8. HWLVM05 R Preferred epi	R	ID NO. 8222 as residues: Asn-17 to Lys-27.
HWLUJ19R ID NO. 8231 as residues: Ser-49 to Ser-55. HWLUL47 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8236 as residues: Lys-18 to Lys-24. HWLUL65 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8237 as residues: Lys-18 to Lys-24. HWLUQ87 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8254 as residues: Cys-34 to Arg-41. HWLUR41 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8256 as residues: Cys-34 to Arg-41. HWLUU88 R ID NO. 8256 as residues: Ser-24 to Trp-30. HWLUU88 R ID NO. 8266 as residues: Pro-9 to Gly-20. HWLUV67 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8266 as residues: Pro-9 to Gly-20. HWLUV07 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8268 as residues: Pro-5 to Arg-13. HWLUZ07 R ID NO. 8269 as residues: Glu-1 to Gly-8. HWLVD26 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8278 as residues: Arg-11 to Asp-16. HWLVD74 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8278 as residues: His-1 to Thr-10. HWLVE21 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8278 as residues: His-1 to Thr-10. HWLVF34 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8282 as residues: Leu-33 to Glu-40, Lys-52 to Lys-62. HWLVF34 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8282 as residues: Arg-54 to His-62. HWLVF34 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8290 as residues: Arg-54 to His-62. HWLV116R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8290 as residues: Arg-1 to Thr-8. HWLVV106 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8293 as residues: Arg-1 to Thr-8. HWLVN05 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8301 as residues: Arg-1 to Arg-14. HWLVW09 Preferred epi	HWLUH72	Preferred epitopes include those comprising a sequence shown in SEQ
HWLUL47 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8236 as residues: Lys-18 to Lys-24. HWLUL65 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8237 as residues: Asp-1 to His-8. HWLUQ87 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8237 as residues: Asp-1 to His-8. HWLUR41 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8254 as residues: Cys-34 to Arg-41. HWLUU88 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8266 as residues: Pro-9 to Gly-20. HWLUV67 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8261 as residues: Pro-9 to Arg-13. HWLUZ07 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8269 as residues: Glu-1 to Gly-8. HWLVD26 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8275 as residues: Arg-11 to Asp-16. HWLVD74 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8278 as residues: His-1 to Thr-10. HWLVE21 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8278 as residues: Leu-33 to Glu-40, Lys-52 to Lys-62. HWLVF34 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8282 as residues: Arg-54 to His-62. HWLVJ15R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8282 as residues: Arg-54 to His-62. HWLVJ84R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8290 as residues: Asr-2 to Gly-33, Ser-35 to Phe-63. HWLVK62 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8298 as residues: Asr-2 to Gly-33, Ser-35 to Phe-63. HWLVM05 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8298 as residues: Ser-1 to Glu-13. HWLVW06 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8301 as residues: Arg-1 to Th-8. HWLVW09 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 83		
HWLUL47 R ID NO. 8236 as residues: Lys-18 to Lys-24. HWLUL65 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8237 as residues: Asp-1 to His-8. HWLUQ87 R ID NO. 8237 as residues: Cys-34 to Arg-41. HWLUR41 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8254 as residues: Cys-34 to Arg-41. HWLUR41 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8256 as residues: Ser-24 to Trp-30. HWLUU88 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8261 as residues: Pro-9 to Gly-20. HWLUV67 R ID NO. 8264 as residues: Pro-9 to Gly-20. HWLUV67 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8264 as residues: Pro-5 to Arg-13. HWLUD20 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8269 as residues: Glu-1 to Gly-8. HWLVD26 R ID NO. 8275 as residues: Arg-11 to Asp-16. HWLVD74 R ID NO. 8278 as residues: His-1 to Thr-10. HWLVE21 R ID NO. 8279 as residues: His-1 to Thr-10. HWLVF34 R ID NO. 8282 as residues: Leu-33 to Glu-40, Lys-52 to Lys-62. HWLV734 R ID NO. 8282 as residues: Arg-54 to His-62. HWLVJ15R ID NO. 8282 as residues: Arg-54 to His-62. HWLVJ84R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8289 as residues: Asp-2 to Gly-33, Scr-35 to Phe-63. HWLVS42 R ID NO. 8290 as residues: Ser-1 to Gly-33, Scr-35 to Phe-63. HWLVK62 R ID NO. 8290 as residues: Ser-1 to Gly-33, Scr-35 to Phe-63. HWLVK62 R ID NO. 8293 as residues: Asp-2 to Thr-8. HWLVM05 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8298 as residues: Asp-1 to Thr-8. HWLVM05 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8298 as residues: Asp-1 to Thr-8. HWLVW06 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8301 as residues: Asp-1 to Thr-8. HWLVW06 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8317 as residues: Gln-1 to Tyr-6. HWLVW09 Preferred epitopes i	HWLUJ19R	
R ID NO. 8236 as residues: Lys-18 to Lys-24. HWLUG55 R ID NO. 8237 as residues: Asp-1 to His-8. HWLUQ87 R ID NO. 8254 as residues: Asp-1 to His-8. HWLUR41 R ID NO. 8256 as residues: Cys-34 to Arg-41. HWLUR41 R ID NO. 8256 as residues: Cys-34 to Arg-41. HWLUU88 R ID NO. 8256 as residues: Ser-24 to Trp-30. HWLUU88 R ID NO. 8261 as residues: Pro-9 to Gly-20. HWLUV67 R ID NO. 8261 as residues: Pro-9 to Gly-20. HWLUV707 R ID NO. 8264 as residues: Pro-9 to Arg-13. HWLUD207 R ID NO. 8264 as residues: Pro-5 to Arg-13. HWLUD207 R ID NO. 8269 as residues: Glu-1 to Gly-8. HWLVD206 R ID NO. 8275 as residues: Arg-11 to Asp-16. HWLVD74 R ID NO. 8275 as residues: His-1 to Thr-10. HWLVD74 R ID NO. 8279 as residues: Leu-33 to Glu-40, Lys-52 to Lys-62. HWLVF34 R ID NO. 8282 as residues: Leu-33 to Glu-40, Lys-52 to Lys-62. HWLVJ158 R ID NO. 8289 as residues: Arg-54 to His-62. HWLVJ158 R ID NO. 8289 as residues: Pro-3 to Gly-33, Ser-35 to Phe-63. HWLVJ84R R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8289 as residues: Pro-3 to Gly-33, Ser-35 to Phe-63. HWLVJ84R R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8289 as residues: Arg-54 to His-62. HWLVJ168 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8289 as residues: Arg-54 to His-62. HWLVJ84R R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8290 as residues: Arg-54 to His-62. HWLVJ84R R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8290 as residues: Arg-54 to His-62. HWLVM62 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8290 as residues: Arg-1 to Thr-8. HWLVM64 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8298 as residues: Arg-1 to Thr-8. HWLVM05 R ID NO. 8301 as residues: Arg-1 to Thr-8. HWLVW06 R ID NO. 83117 as residues: Arg-1 to Arg-14. HWLVW89 R ID NO. 8317 as residues: Arg-1 to Arg-14. HWLVW89 R ID NO. 8317 as residues: Arg-1 to Arg-14.	HWI III 47	
HWLUL65 R ID NO. 8237 as residues: Asp-1 to His-8. HWLUQ87 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8254 as residues: Cys-34 to Arg-41. HWLUR41 R ID NO. 8256 as residues: Ser-24 to Trp-30. HWLUU88 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8256 as residues: Ser-24 to Trp-30. HWLUU88 R ID NO. 8261 as residues: Pro-9 to Gly-20. HWLUV67 R ID NO. 8264 as residues: Pro-9 to Gly-20. HWLUZ07 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8264 as residues: Pro-5 to Arg-13. HWLVD704 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8269 as residues: Glu-1 to Gly-8. HWLVD26 R ID NO. 8278 as residues: Arg-11 to Asp-16. HWLVD74 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8278 as residues: His-1 to Thr-10. HWLVE21 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8278 as residues: His-1 to Thr-10. HWLVF34 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8282 as residues: Leu-33 to Glu-40, Lys-52 to Lys-62. HWLVJ15R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8282 as residues: Arg-54 to His-62. HWLVJ15R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8280 as residues: Arg-54 to His-62. HWLVJ84R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8290 as residues: Asp-2 to Gly-33, Ser-35 to Phe-63. HWLVK62 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8295 as residues: Asp-1 to Thr-8. HWLVM05 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8298 as residues: Asp-1 to Thr-8. HWLVM05 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8298 as residues: Ala-8 to Asn-15. HWLVW06 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8310 as residues: Gln-1 to Tyr-6. HWLVW06 Preferred epitopes include those comprising a sequence shown in S		
R ID NO. 8237 as residues: Asp-1 to His-8. HWLUQ87 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8254 as residues: Cys-34 to Arg-41. HWLUR41 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8256 as residues: Ser-24 to Trp-30. HWLUU88 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8261 as residues: Pro-9 to Gly-20. HWLUV67 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8264 as residues: Pro-5 to Arg-13. HWLUZ07 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8269 as residues: Glu-1 to Gly-8. HWLVD26 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8275 as residues: Arg-11 to Asp-16. HWLVD74 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8278 as residues: His-1 to Thr-10. HWLVE1 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8279 as residues: His-1 to Thr-10. HWLV51 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8282 as residues: Leu-33 to Glu-40, Lys-52 to Lys-62. HWLVJ15R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8282 as residues: Arg-54 to His-62. HWLVJ184R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8290 as residues: Phe-38 to Phe-44. HWLVJ84R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8290 as residues: Asn-2 to Gly-33, Scr-35 to Phe-63. HWLVK62 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8295 as residues: Asn-2 to Gly-33, Scr-35 to Phe-63. HWLVV10 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8295 as residues: Asn-2 to Thr-8. HWLVM05 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8298 as residues: Asn-2 to Thr-8. HWLVW06 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8301 as residues: Asn-6 to Gly-11.		
HWLUQ87 R DNO. 8254 as residues: Cys-34 to Arg-41. HWLUR41 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8256 as residues: Ser-24 to Trp-30. HWLUU88 R DNO. 8261 as residues: Pro-9 to Gly-20. HWLUV67 R ID NO. 8264 as residues: Pro-9 to Gly-20. HWLUZ07 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8264 as residues: Pro-5 to Arg-13. HWLUZ07 R ID NO. 8264 as residues: Glu-1 to Gly-8. HWLVD26 R ID NO. 8275 as residues: Arg-11 to Asp-16. HWLVD74 R ID NO. 8278 as residues: His-1 to Thr-10. HWLVE21 R ID NO. 8279 as residues: Leu-33 to Glu-40, Lys-52 to Lys-62. HWLVF34 R ID NO. 8289 as residues: Arg-54 to His-62. HWLVJ15R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8289 as residues: Arg-54 to His-62. HWLVJ15R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8298 as residues: Arg-54 to His-62. HWLVJ15R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8298 as residues: Arg-54 to His-62. HWLVJ16R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8290 as residues: Arg-54 to His-62. HWLVJ84R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8290 as residues: Arg-54 to His-62. HWLVJ64P Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8290 as residues: Arg-10 to His-62. HWLVJ64P Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8290 as residues: Arg-1 to Thr-8. HWLVL10 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8295 as residues: Arg-1 to Thr-8. HWLVM05 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8298 as residues: Arg-1 to Thr-6. HWLVN12 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8301 as residues: Gln-1 to Tyr-6. HWLVV06 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8310 as residues: Arg-1 to Arg-14. HWLVW89 Preferred epitopes include those compris	1	
R ID NO. 8254 as residues: Cys-34 to Arg-41. HWLUR41 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8256 as residues: Ser-24 to Trp-30. HWLUU88 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8261 as residues: Pro-9 to Gly-20. HWLUV67 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8264 as residues: Pro-5 to Arg-13. HWLUZ07 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8269 as residues: Glu-1 to Gly-8. HWLVD26 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8275 as residues: Arg-11 to Asp-16. HWLVD74 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8278 as residues: His-1 to Thr-10. HWLVE21 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8279 as residues: Leu-33 to Glu-40, Lys-52 to Lys-62. HWLVF34 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8282 as residues: Arg-54 to His-62. HWLVJ15R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8289 as residues: Phe-38 to Phe-44. HWLVJ84R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8290 as residues: Asn-2 to Gly-33, Scr-35 to Phe-63. HWLVK62 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8292 as residues: Asn-2 to Gly-33, Scr-35 to Phe-63. HWLVL10 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8295 as residues: Arg-1 to Glu-13. HWLVN05 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8298 as residues: Arg-1 to Thr-8. HWLVN12 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8301 as residues: Arg-1 to Thr-8. HWLVN14 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8301 as residues: Arg-1 to Tyr-6. HWLVV06 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8310 as residues: Arg-1 to Arg-14.		
HWLUR41 R DNO. 8256 as residues: Ser-24 to Trp-30. HWLUU88 R DNO. 8261 as residues: Pro-9 to Gly-20. HWLUV67 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8261 as residues: Pro-9 to Gly-20. HWLUZ07 R DNO. 8264 as residues: Pro-5 to Arg-13. HWLUZ07 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8269 as residues: Glu-1 to Gly-8. HWLVD26 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8275 as residues: Arg-11 to Asp-16. HWLVD74 R DNO. 8278 as residues: His-1 to Thr-10. HWLV21 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8278 as residues: His-1 to Thr-10. HWLVF34 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8279 as residues: Leu-33 to Glu-40, Lys-52 to Lys-62. HWLVJ15R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8282 as residues: Arg-54 to His-62. HWLVJ44R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8289 as residues: Phe-38 to Phe-44. HWLVJ84R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8290 as residues: Asn-2 to Gly-33, Scr-35 to Phe-63. HWLVK62 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8292 as residues: Asn-2 to Gly-33, Scr-35 to Phe-63. HWLVK62 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8298 as residues: Asn-2 to Thr-8. HWLVM05 R DNO. 8298 as residues: Afg-1 to Thr-8. HWLVM05 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8298 as residues: Afg-1 to Thr-8. HWLVN12 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8301 as residues: Afg-1 to Tyr-6. HWLVV06 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8310 as residues: Afg-1 to Tyr-6. HWLVW09 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8310 as residues: Afg-1 to Afg-14.		
R ID NO. 8256 as residues: Ser-24 to Trp-30. HWLUU88 R ID NO. 8261 as residues: Pro-9 to Gly-20. HWLUV67 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8264 as residues: Pro-5 to Arg-13. HWLUZ07 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8269 as residues: Glu-1 to Gly-8. HWLVD26 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8275 as residues: Arg-11 to Asp-16. HWLVD74 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8278 as residues: His-1 to Thr-10. HWLVE21 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8278 as residues: His-1 to Thr-10. HWLV534 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8282 as residues: Arg-54 to His-62. HWLVJ15R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8282 as residues: Arg-54 to His-62. HWLVJ84R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8280 as residues: Phe-38 to Phe-44. HWLVJ84R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8290 as residues: Asn-2 to Gly-33, Ser-35 to Phe-63. HWLVK62 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8292 as residues: Ser-1 to Glu-13. HWLVU10 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8295 as residues: Arg-1 to Thr-8. HWLVN05 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8298 as residues: Ala-8 to Asn-15. HWLVN02 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8301 as residues: Gln-1 to Tyr-6. HWLVV06 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8310 as residues: Arg-1 to Tyr-6. HWLVV08 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8310 as residues: Arg-1 to Arg-14. HWLVW89 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8317 as residues: Arg-1 to Arg-14.		
HWLUU88 R BY R BY BY BY BY BY BY BY BY BY BY BY BY BY	1	
R ID NO. 8261 as residues: Pro-9 to Gly-20. HWLUV67 R ID NO. 8264 as residues: Pro-5 to Arg-13. HWLUZ07 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8269 as residues: Glu-1 to Gly-8. HWLVD26 R ID NO. 8275 as residues: Arg-11 to Asp-16. HWLVD74 R ID NO. 8278 as residues: His-1 to Thr-10. HWLVD74 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8278 as residues: His-1 to Thr-10. HWLVE21 R ID NO. 8279 as residues: Leu-33 to Glu-40, Lys-52 to Lys-62. HWLVF34 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8279 as residues: Arg-54 to His-62. HWLVJ15R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8289 as residues: Phe-38 to Phe-44. HWLVJ84R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8290 as residues: Phe-38 to Phe-463. HWLVK62 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8290 as residues: Ser-1 to Glu-13. HWLVK62 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8295 as residues: Arg-1 to Thr-8. HWLVM05 R DNO. 8295 as residues: Arg-1 to Thr-8. HWLVN05 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8298 as residues: Ala-8 to Asn-15. HWLVN12 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8298 as residues: Ala-8 to Asn-15. HWLVN10 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8301 as residues: Gln-1 to Tyr-6. HWLVV06 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8310 as residues: Arg-1 to Arg-14. HWLVW89 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8317 as residues: Asn-6 to Gly-11.	HWLUU88	
HWLUV67 R ID NO. 8264 as residues: Pro-5 to Arg-13. HWLUZ07 R ID NO. 8269 as residues: Glu-1 to Gly-8. HWLVD26 R ID NO. 8269 as residues: Glu-1 to Gly-8. HWLVD26 R ID NO. 8275 as residues: Glu-1 to Asp-16. HWLVD74 R ID NO. 8278 as residues: His-1 to Thr-10. HWLVE21 R ID NO. 8278 as residues: His-1 to Thr-10. HWLVF34 R ID NO. 8279 as residues: Leu-33 to Glu-40, Lys-52 to Lys-62. HWLVJ15R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8282 as residues: Arg-54 to His-62. HWLVJ15R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8289 as residues: Phe-38 to Phe-44. HWLVJ84R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8290 as residues: Phe-38 to Phe-44. HWLVK62 R HWLVK62 R HWLVK62 R HWLVK64 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8292 as residues: Ser-1 to Glu-13. HWLVL10 R HWLVL10 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8295 as residues: Ser-1 to Thr-8. HWLVM05 R ID NO. 8298 as residues: Arg-1 to Thr-8. HWLVN12 R HWLVN12 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8298 as residues: Ala-8 to Asn-15. HWLVN06 R ID NO. 8301 as residues: Gln-1 to Tyr-6. HWLVV06 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8310 as residues: Arg-1 to Arg-14. HWLVW89 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8310 as residues: Arg-1 to Arg-14. HWLVW89 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8310 as residues: Arg-1 to Arg-14.	1	
HWLUZ07 R ID NO. 8269 as residues: Glu-1 to Gly-8. HWLVD26 R ID NO. 8275 as residues: Arg-11 to Asp-16. HWLVD74 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8275 as residues: Arg-11 to Asp-16. HWLVD74 R ID NO. 8278 as residues: His-1 to Thr-10. HWLVE21 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8279 as residues: Leu-33 to Glu-40, Lys-52 to Lys-62. HWLVF34 R ID NO. 8282 as residues: Arg-54 to His-62. HWLVJ15R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8289 as residues: Phe-38 to Phe-44. HWLVJ84R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8290 as residues: Asn-2 to Gly-33, Ser-35 to Phe-63. HWLVK62 R ID NO. 8290 as residues: Ser-1 to Glu-13. HWLVL10 R ID NO. 8292 as residues: Arg-1 to Thr-8. HWLVM05 R ID NO. 8298 as residues: Arg-1 to Thr-8. HWLVM05 R ID NO. 8298 as residues: Ala-8 to Asn-15. HWLVN12 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8301 as residues: Gln-1 to Tyr-6. HWLVV06 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8310 as residues: Arg-1 to Arg-14. HWLVW89 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8310 as residues: Arg-1 to Arg-14.	HWLUV67	Preferred epitopes include those comprising a sequence shown in SEQ
R ID NO. 8269 as residues: Glu-1 to Gly-8. HWLVD26 R ID NO. 8275 as residues: Arg-11 to Asp-16. HWLVD74 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8278 as residues: His-1 to Thr-10. HWLVE21 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8279 as residues: Leu-33 to Glu-40, Lys-52 to Lys-62. HWLVF34 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8282 as residues: Arg-54 to His-62. HWLVJ15R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8289 as residues: Phe-38 to Phe-44. HWLVJ84R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8290 as residues: Asn-2 to Gly-33, Scr-35 to Phe-63. HWLVK62 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8292 as residues: Ser-1 to Glu-13. HWLVL10 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8295 as residues: Arg-1 to Thr-8. HWLVM05 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8298 as residues: Ala-8 to Asn-15. HWLVN12 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8301 as residues: Ala-8 to Asn-15. HWLVV06 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8301 as residues: Ala-8 to Arg-1 to Arg-14. HWLVW89 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8310 as residues: Arg-1 to Arg-14.	R	ID NO. 8264 as residues: Pro-5 to Arg-13.
HWLVD26 R ID NO. 8275 as residues: Arg-11 to Asp-16. HWLVD74 R ID NO. 8278 as residues: Arg-11 to Asp-16. HWLVE21 R ID NO. 8278 as residues: His-1 to Thr-10. HWLVE21 R ID NO. 8279 as residues: His-1 to Thr-10. HWLVF34 R ID NO. 8279 as residues: Leu-33 to Glu-40, Lys-52 to Lys-62. HWLVF34 R ID NO. 8282 as residues: Arg-54 to His-62. HWLVJ15R ID NO. 8282 as residues: Arg-54 to His-62. HWLVJ44R HWLVJ45R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8289 as residues: Phe-38 to Phe-44. HWLVJ84R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8290 as residues: Asn-2 to Gly-33, Ser-35 to Phe-63. HWLVK62 R ID NO. 8292 as residues: Ser-1 to Glu-13. HWLVL10 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8295 as residues: Arg-1 to Thr-8. HWLVM05 R ID NO. 8298 as residues: Ala-8 to Asn-15. HWLVN12 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8301 as residues: Gln-1 to Tyr-6. HWLVV06 R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8310 as residues: Arg-1 to Arg-14. HWLVW89 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8317 as residues: Asn-6 to Gly-11.	HWLUZ07	Preferred epitopes include those comprising a sequence shown in SEQ
R ID NO. 8275 as residues: Arg-11 to Asp-16. HWLVD74 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8278 as residues: His-1 to Thr-10. HWLVE21 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8279 as residues: Leu-33 to Glu-40, Lys-52 to Lys-62. HWLVF34 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8282 as residues: Arg-54 to His-62. HWLVJ15R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8289 as residues: Phe-38 to Phe-44. HWLVJ84R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8290 as residues: Asn-2 to Gly-33, Ser-35 to Phe-63. HWLVK62 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8292 as residues: Ser-1 to Glu-13. HWLVL10 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8295 as residues: Arg-1 to Thr-8. HWLVM05 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8298 as residues: Ala-8 to Asn-15. HWLVN12 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8301 as residues: Gln-1 to Tyr-6. HWLVV06 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8310 as residues: Arg-1 to Arg-14. HWLVW89 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8317 as residues: Asn-6 to Gly-11.	R	ID NO. 8269 as residues: Glu-1 to Gly-8.
HWLVD74 R ID NO. 8278 as residues: His-1 to Thr-10. HWLVE21 R ID NO. 8279 as residues: His-1 to Thr-10. HWLVF34 R ID NO. 8282 as residues: Leu-33 to Glu-40, Lys-52 to Lys-62. HWLVF34 R ID NO. 8282 as residues: Arg-54 to His-62. HWLVJ15R ID NO. 8289 as residues: Phe-38 to Phe-44. HWLVJ84R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8290 as residues: Asn-2 to Gly-33, Ser-35 to Phe-63. HWLVK62 R ID NO. 8292 as residues: Ser-1 to Glu-13. HWLVL10 R ID NO. 8295 as residues: Arg-1 to Thr-8. HWLVM05 R ID NO. 8298 as residues: Ala-8 to Asn-15. HWLVN12 R ID NO. 8301 as residues: Gln-1 to Tyr-6. HWLVV06 R ID NO. 8310 as residues: Arg-1 to Arg-14. HWLVW89 R ID NO. 8317 as residues: Asn-6 to Gly-11.	HWLVD26	Preferred epitopes include those comprising a sequence shown in SEQ
R ID NO. 8278 as residues: His-1 to Thr-10. HWLVE21 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8279 as residues: Leu-33 to Glu-40, Lys-52 to Lys-62. HWLVF34 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8282 as residues: Arg-54 to His-62. HWLVJ15R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8289 as residues: Phe-38 to Phe-44. HWLVJ84R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8290 as residues: Asn-2 to Gly-33, Ser-35 to Phe-63. HWLVK62 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8292 as residues: Ser-1 to Glu-13. HWLVL10 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8295 as residues: Arg-1 to Thr-8. HWLVM05 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8298 as residues: Ala-8 to Asn-15. HWLVN12 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8301 as residues: Gln-1 to Tyr-6. HWLVV06 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8310 as residues: Arg-1 to Arg-14. HWLVW89 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8317 as residues: Asn-6 to Gly-11.	R	ID NO. 8275 as residues: Arg-11 to Asp-16.
HWLVE21 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8279 as residues: Leu-33 to Glu-40, Lys-52 to Lys-62. HWLVF34 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8282 as residues: Arg-54 to His-62. HWLVJ15R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8289 as residues: Phe-38 to Phe-44. HWLVJ84R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8290 as residues: Asn-2 to Gly-33, Scr-35 to Phe-63. HWLVK62 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8292 as residues: Ser-1 to Glu-13. HWLVL10 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8295 as residues: Arg-1 to Thr-8. HWLVM05 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8298 as residues: Ala-8 to Asn-15. HWLVN12 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8301 as residues: Gln-1 to Tyr-6. HWLVV06 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8310 as residues: Arg-1 to Arg-14. HWLVW89 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8317 as residues: Asn-6 to Gly-11.	HWLVD74	
R ID NO. 8279 as residues: Leu-33 to Glu-40, Lys-52 to Lys-62. HWLVF34 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8282 as residues: Arg-54 to His-62. HWLVJ15R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8289 as residues: Phe-38 to Phe-44. HWLVJ84R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8290 as residues: Asn-2 to Gly-33, Ser-35 to Phe-63. HWLVK62 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8292 as residues: Ser-1 to Glu-13. HWLVL10 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8295 as residues: Arg-1 to Thr-8. HWLVM05 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8298 as residues: Ala-8 to Asn-15. HWLVN12 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8301 as residues: Gln-1 to Tyr-6. HWLVV06 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8310 as residues: Arg-1 to Arg-14. HWLVW89 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8317 as residues: Asn-6 to Gly-11.		
HWLVF34 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8282 as residues: Arg-54 to His-62. HWLVJ15R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8289 as residues: Phe-38 to Phe-44. HWLVJ84R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8290 as residues: Asn-2 to Gly-33, Ser-35 to Phe-63. HWLVK62 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8292 as residues: Ser-1 to Glu-13. HWLVL10 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8295 as residues: Arg-1 to Thr-8. HWLVM05 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8298 as residues: Ala-8 to Asn-15. HWLVN12 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8301 as residues: Gln-1 to Tyr-6. HWLVV06 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8310 as residues: Arg-1 to Arg-14. HWLVW89 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8317 as residues: Asn-6 to Gly-11.		
R ID NO. 8282 as residues: Arg-54 to His-62. HWLVJ15R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8289 as residues: Phe-38 to Phe-44. HWLVJ84R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8290 as residues: Asn-2 to Gly-33, Ser-35 to Phe-63. HWLVK62 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8292 as residues: Ser-1 to Glu-13. HWLVL10 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8295 as residues: Arg-1 to Thr-8. HWLVM05 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8298 as residues: Ala-8 to Asn-15. HWLVN12 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8301 as residues: Gln-1 to Tyr-6. HWLVV06 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8310 as residues: Arg-1 to Arg-14. HWLVW89 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8317 as residues: Asn-6 to Gly-11.		
HWLVJ84R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8289 as residues: Phe-38 to Phe-44. HWLVJ84R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8290 as residues: Asn-2 to Gly-33, Ser-35 to Phe-63. HWLVK62 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8292 as residues: Ser-1 to Glu-13. HWLVL10 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8295 as residues: Arg-1 to Thr-8. HWLVM05 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8298 as residues: Ala-8 to Asn-15. HWLVN12 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8301 as residues: Gln-1 to Tyr-6. HWLVV06 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8310 as residues: Arg-1 to Arg-14. HWLVW89 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8317 as residues: Asn-6 to Gly-11.		
ID NO. 8289 as residues: Phe-38 to Phe-44. HWLVJ84R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8290 as residues: Asn-2 to Gly-33, Ser-35 to Phe-63. HWLVK62 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8292 as residues: Ser-1 to Glu-13. HWLVL10 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8295 as residues: Arg-1 to Thr-8. HWLVM05 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8298 as residues: Ala-8 to Asn-15. HWLVN12 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8301 as residues: Gln-1 to Tyr-6. HWLVV06 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8310 as residues: Arg-1 to Arg-14. HWLVW89 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8317 as residues: Asn-6 to Gly-11.		
HWLVJ84R Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8290 as residues: Asn-2 to Gly-33, Ser-35 to Phe-63. HWLVK62 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8292 as residues: Ser-1 to Glu-13. HWLVL10 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8295 as residues: Arg-1 to Thr-8. HWLVM05 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8298 as residues: Ala-8 to Asn-15. HWLVN12 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8301 as residues: Gln-1 to Tyr-6. HWLVV06 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8310 as residues: Arg-1 to Arg-14. HWLVW89 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8317 as residues: Asn-6 to Gly-11.	HWLVIISK	,
ID NO. 8290 as residues: Asn-2 to Gly-33, Ser-35 to Phe-63. HWLVK62 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8292 as residues: Ser-1 to Glu-13. HWLVL10 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8295 as residues: Arg-1 to Thr-8. HWLVM05 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8298 as residues: Ala-8 to Asn-15. HWLVN12 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8301 as residues: Gln-1 to Tyr-6. HWLVV06 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8310 as residues: Arg-1 to Arg-14. HWLVW89 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8317 as residues: Asn-6 to Gly-11.	HWI WIGAD	
HWLVK62 R D NO. 8292 as residues: Ser-1 to Glu-13. HWLVL10 R D NO. 8295 as residues: Ser-1 to Thr-8. HWLVM05 R D NO. 8298 as residues: Arg-1 to Thr-8. HWLVN12 R D NO. 8298 as residues: Ala-8 to Asn-15. HWLVN12 R D NO. 8301 as residues: Gln-1 to Tyr-6. HWLVV06 R D NO. 8310 as residues: Arg-1 to Arg-14. HWLVW89 R D NO. 8317 as residues: Asn-6 to Gly-11.	I TWLVJ84K	
R ID NO. 8292 as residues: Ser-1 to Glu-13. HWLVL10 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8295 as residues: Arg-1 to Thr-8. HWLVM05 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8298 as residues: Ala-8 to Asn-15. HWLVN12 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8301 as residues: Gln-1 to Tyr-6. HWLVV06 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8310 as residues: Arg-1 to Arg-14. HWLVW89 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8317 as residues: Asn-6 to Gly-11.	HWI VK62	
HWLVL10 R ID NO. 8295 as residues: Arg-1 to Thr-8. HWLVM05 R ID NO. 8298 as residues: Ala-8 to Asn-15. HWLVN12 R ID NO. 8301 as residues: Gln-1 to Tyr-6. HWLVV06 R ID NO. 8310 as residues: Arg-1 to Arg-14. HWLVW89 R ID NO. 8317 as residues: Asn-6 to Gly-11.	1	
R ID NO. 8295 as residues: Arg-1 to Thr-8. HWLVM05 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8298 as residues: Ala-8 to Asn-15. HWLVN12 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8301 as residues: Gln-1 to Tyr-6. HWLVV06 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8310 as residues: Arg-1 to Arg-14. HWLVW89 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8317 as residues: Asn-6 to Gly-11.	<u> </u>	
HWLVM05 R D NO. 8298 as residues: Ala-8 to Asn-15. HWLVN12 R D NO. 8301 as residues: Gln-1 to Tyr-6. HWLVV06 R D NO. 8310 as residues: Arg-1 to Arg-14. HWLVW89 R D NO. 8317 as residues: Asn-6 to Gly-11.		
R ID NO. 8298 as residues: Ala-8 to Asn-15. HWLVN12 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8301 as residues: Gln-1 to Tyr-6. HWLVV06 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8310 as residues: Arg-1 to Arg-14. HWLVW89 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8317 as residues: Asn-6 to Gly-11.		
HWLVN12 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8301 as residues: Gln-1 to Tyr-6. HWLVV06 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8310 as residues: Arg-1 to Arg-14. HWLVW89 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8317 as residues: Asn-6 to Gly-11.		
R ID NO. 8301 as residues: Gln-1 to Tyr-6. HWLVV06 R ID NO. 8310 as residues: Arg-1 to Arg-14. HWLVW89 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8317 as residues: Asn-6 to Gly-11.		
HWLVV06 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8310 as residues: Arg-1 to Arg-14. HWLVW89 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8317 as residues: Asn-6 to Gly-11.		
R ID NO. 8310 as residues: Arg-1 to Arg-14. HWLVW89 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8317 as residues: Asn-6 to Gly-11.		
HWLVW89 Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8317 as residues: Asn-6 to Gly-11.		
R ID NO. 8317 as residues: Asn-6 to Gly-11.		
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HWLVY14 Preferred epitopes include those comprising a sequence shown in SEQ	HWLVY14	Preferred epitopes include those comprising a sequence shown in SEQ

R	ID NO. 8320 as residues: Ser-1 to Trp-7.
HWLWA14	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8324 as residues: Thr-1 to Trp-9.
HWLWA82	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8325 as residues: Val-1 to Ser-8, Arg-52 to Gly-58.
HWLWB71	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8331 as residues: Cys-28 to Trp-42.
HWLWB77	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8333 as residues: Cys-40 to Trp-47.
HWLWD32	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8334 as residues: Gly-13 to Ala-21.
HWLWD60	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8336 as residues: Tyr-16 to Phe-22.
HWLWE80	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8339 as residues: Gly-1 to Trp-6.
HWLWJ36	Preferred epitopes include those comprising a sequence shown in SEQ
· R	ID NO. 8346 as residues: Asp-11 to Asn-25.
HWLWO57	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8353 as residues: Ser-1 to Phe-6.
HWLWP08	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8357 as residues: Arg-4 to Val-12.
HWLWS28	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8369 as residues: Arg-4 to Tyr-9.
HWLWU27	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8373 as residues: Ala-16 to Phe-21.
HWLWW4	Preferred epitopes include those comprising a sequence shown in SEQ
6R	ID NO. 8374 as residues: Ser-6 to Ser-16.
HWLXA13	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8380 as residues: Asp-8 to Ser-17.
HWLXA23	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8381 as residues: Pro-10 to Ile-20.
HWLXJ59R	Preferred epitopes include those comprising a sequence shown in SEQ
171117 373 700	ID NO. 8388 as residues: Pro-7 to Ser-13.
HWLXN33	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8391 as residues: Glu-1 to Gly-7.
HWLXP33	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8395 as residues: Thr-3 to Lys-13.
HWLXP45	Preferred epitopes include those comprising a sequence shown in SEQ ID NO. 8396 as residues: Gly-10 to Gly-22, Pro-27 to Arg-35.
R HWLXR49	Preferred epitopes include those comprising a sequence shown in SEQ
	ID NO. 8403 as residues: Gly-10 to Pro-15.
R HWLXT31	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8405 as residues: Gly-10 to Glu-15, Ser-31 to Lys-36.
HWMBC46	Preferred epitopes include those comprising a sequence shown in SEQ
R R	ID NO. 8410 as residues: Phe-11 to Lys-17.
HWMBD22	Preferred epitopes include those comprising a sequence shown in SEQ
R R	ID NO. 8411 as residues: Pro-11 to Ala-18.
HWMBD71	Preferred epitopes include those comprising a sequence shown in SEQ
R R	ID NO. 8413 as residues: Asp-4 to Leu-9.
	1 D 110. 0713 as residues. Asp-1 to Lea-7.

HWMBE36	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8415 as residues: Tyr-12 to Met-18.
HWMBF87	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8416 as residues: Gly-1 to Arg-6.
HWMBG63	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8417 as residues: Glu-1 to Ser-9.
HWMBI08	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8420 as residues: Arg-29 to His-37, Trp-43 to Arg-48.
HWMBK47	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8423 as residues: Asn-6 to His-11, Asn-25 to Cys-30.
HWMBL29	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8425 as residues: Leu-11 to Phe-16.
HWMBL57	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8426 as residues: Glu-46 to Tyr-57.
HWMBL82	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8427 as residues: Leu-27 to Thr-67, His-74 to Asn-79, Ser-83 to
	Lys-94, Gln-109 to Lys-115, Asp-122 to Tyr-131, Leu-138 to Arg-145,
	Glu-149 to Lys-154.
HWMBM67	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8430 as residues: Gly-32 to Arg-37, Ala-41 to Asp-47.
HWMBM83	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8431 as residues: Thr-21 to Asn-31.
HWMBN52	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8435 as residues: Thr-21 to Glu-34, Leu-50 to Cys-56.
HWMBP01	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8437 as residues: Ser-9 to Ile-17.
HWMBR40	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8443 as residues: Pro-8 to Glu-19.
HWMBR68	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8446 as residues: Tyr-1 to Trp-6.
HWMBR77	Preferred epitopes include those comprising a sequence shown in SEQ
RA	ID NO. 8448 as residues: Lys-1 to Val-8.
HWMBT23	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8454 as residues: Val-3 to Arg-14.
HWMBV48	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8458 as residues: Pro-51 to Ser-57, Gln-65 to Leu-76.
HWMBW5	Preferred epitopes include those comprising a sequence shown in SEQ
4R	ID NO. 8460 as residues: Pro-55 to Glu-63.
HWMBY90	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8466 as residues: Thr-1 to Arg-6.
HWMCB93	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8473 as residues: His-1 to Ala-13.
HWMCE24	Preferred epitopes include those comprising a sequence shown in SEQ
R R	ID NO. 8480 as residues: Asn-6 to Lys-12.
HWMCF45	Preferred epitopes include those comprising a sequence shown in SEQ
R	ID NO. 8482 as residues: Pro-10 to Phe-16.
HWMCH47	Preferred epitopes include those comprising a sequence shown in SEQ
R R	ID NO. 8484 as residues: Pro-16 to Ser-24.
HWMCH76	
11 WIVICITIO	Preferred epitopes include those comprising a sequence shown in SEQ